

## (12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization International Bureau



(43) International Publication Date  
22 April 2004 (22.04.2004)

PCT

(10) International Publication Number  
**WO 2004/032957 A1**

(51) International Patent Classification<sup>7</sup>: **A61K 39/09**, C07K 14/315, 16/12

(74) Agents: ZACCO SWEDEN AB et al.; P.O. Box 23101, Sveavägen 170, S-104 35 Stockholm (SE).

(21) International Application Number: **PCT/SE2003/001587**

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.

(22) International Filing Date: 10 October 2003 (10.10.2003)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:  
60/417,660 11 October 2002 (11.10.2002) US

(84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

(71) Applicants and

(72) Inventors: GUSS, Bengt [SE/SE]; Dag Hammarskjölds väg 238 B, S-756 52 Uppsala (SE). FLOCK, Jan-Ingmar [SE/SE]; Sångarvägen 2, S-167 67 Bromma (SE). JACOBSSON, Karin [SE/SE]; Villavägen 22 Ärentuna, S-743 91 Storvreta (SE). JANZON, Kenneth [SE/SE]; Hägerstensvägen 265, S-129 38 Hägersten (SE). FRYKBERG, Lars [SE/SE]; Villavägen 22, S-743 91 Storvreta (SE). FLOCK, Margareta [SE/SE]; Sångarvägen 2, S-167 67 Bromma (SE). BERGMAN, Rune [SE/SE]; Laxvägen 3, S-147 43 Tumba (SE).

## Published:

— with international search report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

**WO 2004/032957 A1**

(54) Title: IMMUNIZATION OF NON-HUMAN MAMMALS AGAINST STREPTOCOCCUS EQUI

(57) Abstract: The present invention is concerned with an antigenic composition comprising at least one antigen that comprises at least one antigenic epitope or antigenic determinant derived from a protein present in one or both of *S. equi* subsp. *Equi* and subsp. *Zooepidemicus* and use thereof for immunization of non-human mammals against *S. equi* subsp. *equi* and/or subsp. *Zooepidemicus*. The present invention also discloses a vaccine composition comprising the aforesaid antigenic composition as immunizing component. The antigens used are EAG, FNZ, SFS, SEC and ScLC.

IMMUNIZATION OF NON-HUMAN MAMMALS AGAINST *STREPTOCOCCUS EQUI*

5

**Background of the Invention****1. Field of the Invention**

This invention is generally related to antigenic compositions and use thereof for immunization of non-human mammals, e.g. horses, against *Streptococcus equi*.

10 Streptococcal infections in horses are mainly caused by the species *Streptococcus equi*, which is classified as a Lancefield Group C *Streptococcus* and comprises two subspecies designated *equi* and *zooepidemicus*, respectively.

15 *Streptococcus equi* subsp. *equi*, which is virtually confined to horses, is the causative agent of strangles, a world-wide distributed and highly contagious serious disease of the upper respiratory tract of the Equidae. Strangles is one of the most frequently reported equine diseases world-wide and is characterized by fever, nasal discharge, and abscess formation in the retropharyngeal and mandibular lymph nodes. In some cases the disease shows a metastatic course in the body, so called "bastard strangles". The disease has a world-wide distribution and causes great economic losses. Moreover, since strangles is a highly 20 contagious disease, not only infected animals but also all other members of e.g. an afflicted stud must be isolated for as long as up to three months.

25 *S. equi* subsp. *zooepidemicus* is considered as an opportunistic commensal often occurring in the upper respiratory tract of healthy horses. However, after stress or virus infection, it can cause a secondary infection, which results in strangles-like symptoms.

30 Moreover, subsp. *zooepidemicus* infects not only horses but also a wide range of other animals, like pigs, dogs, cats, and cows. Even human cases of infection due to subsp. *zooepidemicus* have been reported. This subspecies has been implicated as the primary pathogen in conditions such as endometritis, cervicitis, abortion, mastitis, pneumonia, abscesses and joint infections.

35 Although it is possible to treat and cure these streptococcal infections with antibiotics, such as penicillin, tetracycline or gentamicin, an effective prophylactic agent that could prevent outbursts of such infections and obviate or reduce the risk for development of resistant strains associated with antibiotic treatment, would be appreciated.

## 2. Description of the Related Art

However, although many attempts have been made to develop prophylactic agents such as vaccines against *S. equi*, at the present time no efficient vaccines or immunizing preparations are available, neither for the subspecies *equi* nor for the subspecies 5 *zooepidemicus*.

Existing vaccines against strangles are based on inactivated, e. g. heat-killed, or attenuated strains of *S. equi* subsp. *equi* or acid extracts/mutanolysin enriched in M-protein(s), i.e. immunogenic protein(s) produced by *S. equi*. A vaccine against *S. equi* subsp. *zooepidemicus* based on an M-like protein is disclosed in US-A-5,583,014. In WO 87/00436, 10 an avirulent strain of *S. equi* is disclosed for use as a vaccine against *S. equi* that stimulates an antibody response in the nasopharyngeal mucosa after administration thereof to a horse.

Since the previously developed vaccines or immunizing preparations are hampered by side-effects and, moreover, provide insufficient protection, there is a need for efficient and safe prophylactic agents, such as vaccines, that protect against *S. equi* infections and/or 15 prevent spread thereof without giving rise to undesirable side-effects.

It is well known that attachment to eukaryotic cell surfaces is an essential step in the establishment of infection and colonization by bacterial pathogens. Accordingly, streptococcal surface proteins, that interact with and/or bind to different components of the Extracellular Matrix (ECM) or plasma proteins of the host cell, are potential candidates for use as active 20 component(s) for immunizing purposes.

This is illustrated by the vaccines based on M-like proteins mentioned above or disclosed in the literature, i.a. in WO 98/0561. The binding of fibrinogen and complement factor H to M-proteins is assumed to be important for the ability of streptococci to resist phagocytosis by polymorphonuclear leucocytes.

25 Another mechanism used by streptococci for attachment to host cells involves binding to the ECM component fibronectin (Fn) (Ref. 21, 22). Binding between Fn-binding bacterial cell-surface proteins and immobilized Fn promotes internalization of streptococci by epithelial cells (Ref. 2, 23, 24). Fibronectin is a dimeric glycoprotein found both in plasma and in a fibrillar form in the extracellular matrix. The main function of Fn is to mediate 30 substrate adhesion of eukaryotic cells, which involves the binding of specific cell-surface receptors to certain domains of the Fn molecule. Furthermore, it also interacts with several other macromolecules, such as DNA, heparin, fibrin, and collagen.

Accordingly, Fn-binding proteins from different streptococcal species have been cloned and sequenced previously. For instance, from *S. equi*, one Fn-binding protein has been

cloned and characterized, which is a Fn-binding cell-surface protein of subsp. *zooepidemicus*, that has been designated FNZ (Lindmark et al., 1996, Ref. 9). Another Fn-binding protein from *S. equi* subsp. *equi*, has been cloned and characterized by Lindmark and Guss (1999) (Ref. 12). This latter protein that is designated SFS and its potential use as an active 5 component in a vaccine for protection of horses against strangles are disclosed in WO 00/37496.

In Jonsson et al. (1995) (Ref. 8), a protein designated ZAG has been cloned and characterized from *S. equi* subsp. *zooepidemicus* that mediates binding to the plasma proteinase inhibitor  $\alpha_2$ M. It is speculated therein that this protein is similar in function to 10 streptococcal M proteins. This protein, ZAG, is also disclosed in WO 95/07296, where its  $\alpha_2$ M-binding properties are indicated. However, immunogenic properties or potential use thereof as an active component in a vaccine for protection of e.g. horses against strangles are not disclosed therein. The gene *zag* encoding ZAG is also disclosed in these references.

A gene that is similar to the aforesaid *zag* gene from *S. equi* subsp. *zooepidemicus* 15 but is present in subsp. *equi* has been described by Lindmark et al. (1999) (Ref. 11) and Lindmark (1999) (Ref. 13). This gene is hereafter designated *eag* and encodes a protein designated EAG.

#### **Brief Summary of the Invention**

The present invention is based on an antigenic composition comprising at least one 20 antigen that comprises at least one antigenic epitope or antigenic determinant derived from a protein present in one or both of *S. equi* subsp. *equi* and subsp. *zooepidemicus* and use thereof for immunization of non-human mammals against *S. equi* subsp. *equi* and/or subsp. *zooepidemicus*.

The present invention is also directed to a vaccine composition comprising the afore- 25 said antigenic composition as immunizing component, methods to prepare said antigenic or vaccine composition, methods to induce an immune response against *S. equi* in non-human mammals and methods for prophylactic or therapeutic treatment of *S. equi* infection in non-human mammals. When used generally, the expression “*S. equi*” refers to one or both of subsp. *equi* and subsp. *zooepidemicus*.

According to a suitable embodiment, the present invention is directed to a vaccine 30 that protects equines, such as horses, against strangles.

In the context of infections caused by *S. equi* subsp. *equi*, the expression “non-human mammals” primarily refers to animals belonging to the family Equidae that consists of horses, donkeys and zebras and to hybrids thereof, such as mules and hinnies.

In connection with infections caused by *S. equi* subsp. *zooepidemicus*, the expression "non-human mammals" in addition refers also to other mammals such as cows, pigs, dogs and cats.

#### **Brief Description of the Drawings**

5 In the following, the present invention is described in closer detail with reference to the drawings, where

**Fig. 1** shows IgG antibodies developed against FNZN in eight individual mice after intranasal inoculation with *Streptococcus equi* subsp. *equi*.

10 **Fig. 2** shows IgG antibodies developed against SFSC1 in eight individual mice after intranasal inoculation with *Streptococcus equi* subsp. *equi*.

**Fig. 3** shows IgG antibodies developed against EAG4B in eight individual mice after intranasal inoculation with *Streptococcus equi* subsp. *equi*.

15 **Fig. 4** shows IgG antibodies developed in mice against FNZN after subcutaneous immunization with FNZN, SFSC1, and EAG4B. Results obtained for seven immunized mice and one non-immunized (▼) mouse are shown.

**Fig. 5** shows IgG antibodies developed in mice against SFSC1 after subcutaneous immunization with FNZN, SFSC1, and EAG4B. Results obtained for seven immunized animals and one non-immunized animal (▼) are shown.

20 **Fig. 6** shows IgG antibodies developed in mice against EAG4B after subcutaneous immunization with FNZN, SFSC1, and EAG4B. Results obtained for seven immunized animals and one non-immunized animal (▼) are shown.

25 **Fig. 7** shows IgG antibodies developed in mice after subcutaneous immunization with SEC 2.16. Results from five immunized and five non-immunized mice are shown. (Pre-immune sera gave no response, values closest to the base line with symbols overlapping each other).

The results shown in **Fig. 1-7**, have been obtained with an ELISA test.

**Fig. 8** shows the accumulated number of mice that survived or lost less than 15% of weight after subcutaneous immunization with FNZN, SFSC1, and EAG4B followed by challenge with *Streptococcus equi* subsp. *equi* (n= 24).

30 **Fig. 9** shows nasal growth after subcutaneous immunization of mice with FNZN, SFSC1, and EAG4B followed by challenge with *Streptococcus equi* subsp. *equi*. Growth is determined on a scale of 0-3 (n= 24). Mean and SE (Standard Error) are shown. The control group was not immunized.

**Fig. 10** shows weight loss of mice after intranasal immunization of mice with FNZN, SFSC1, and EAG4B followed by challenge with *Streptococcus equi* subsp. *equi* (n= 24). The control group was not immunized.

5 **Fig. 11** shows nasal growth after intranasal immunization of mice with FNZN, SFSC1, and EAG4B followed by challenge with *Streptococcus equi* subsp. *equi* (n= 24). Mean and SE are shown.

**Fig. 12** shows IgA antibodies against SFSC1 in bronchoalveolar lavage (BAL) and nasal washes (NW) from mice nasally immunized with FNZN, SFSC1, and EAG4B. Results from six animals are shown.

10 **Fig. 13** shows weight loss of mice after intranasal immunization of mice with EAG4B followed by challenge with *Streptococcus equi* subsp. *equi* (n=10). The control group was not immunized.

**Fig. 14** shows IgG antibodies in immunized horses developed against EAG4B and FNZN.

15 **Fig. 15** shows the results from a comparison of serum antibody titers from horses with or without a history of strangles.

**Fig. 16** shows the presence of IgA antibodies in nasal washings from immunized horses.

20 **Fig. 17** shows antibody titers against EAG4B in horses immunized with EAG4B, FNZN, and SFSC1 as a function of time

**Fig. 18** shows the results obtained for mice (n=15) that were immunized with SEC2.16 using Matrix as adjuvant. The number of animals that lost more than 15% of weight or that died is shown as a function of time.

25 **Fig. 19** shows the results obtained for mice that were immunized with SEC2.16 alone (group B (n=5)) or SEC2.16 together with Matrix (group A (n=5)). Serum samples were taken before (denoted pre) and after (denoted imm) immunization.

#### **Detailed Description of the Invention**

30 The present invention is directed to an antigenic composition comprising at least one antigen, wherein said at least one antigen comprises at least part of a protein of *Streptococcus equi* subsp. *equi*, and said at least part of said protein comprises at least one antigenic epitope or antigenic determinant of *Streptococcus equi*, and wherein said protein is comprised of a

protein that is designated EAG and comprises an N-terminal amino acid sequence, which is shown below as SEQ. ID. NO: 1 and is designated EAG4B.

SEQ. ID. NO: 1:

1 M A L D A T T V L E P T T A F I R E A V R E I N Q L S D D Y A D N Q E L Q A V L A N A G V E A L A A D T V D Q A K A A L  
 5 6 1 D K A K A A V A G V Q L D E A R R E A Y R T I N A L S D Q H K S D Q K V Q L A L V A A A A K V A D A A S V D Q V N A A I  
 12 1 N D A H T A I A D I T G A A L L E A K E A A I N E L K Q Y G I S D Y Y V T L I N K A K T V E G V N A L K A K I L S A L P

According to a suitable embodiment of the present invention, said antigenic composition comprises at least one further antigen that comprises at least part of a fibronectin-binding protein of *Streptococcus equi*, said at least part of said protein comprising at least one 10 antigenic epitope or antigenic determinant of *Streptococcus equi*, and wherein said protein is selected from the group consisting of FNZ (EMBL sequence data bank accession number X99995) comprising an amino acid sequence as shown in SEQ. ID. NO: 2 below:

M K T K S F R K V L T T S A T C I V L A T S F A G G T L R V W A E Q L Y Y G W N D G T R Q S S P Y F L Y V S P K N A P K R E  
 L K D E Y V V Y C F N K K L Y W P D Q W E S I Y S N F N D I R S P Y N D L P V Y E K K L G Y D G I F K Q Y A P D Y K K D I S  
 D I A S A L V A V L S N G Y P T N K S Q L S T S Y H L N N D S S R K V T Q L A I W Y F S D S L T K E Y L K D T G G Y N L N D  
 M E K K A L D F L I S K G E D S K L K S E Q S N Y S L D I Y V Y Q S G G H D H M K D Y Q N L L G S T L I P K E P L K P Q L G  
 G F S G H N G N G L S G L E G G S S G S Q E T N E D G K K G L I G F H G G L S G S E G K R D P L P G L K G E A G A P D T P Q  
 K P N D P L Q G L E G G N S P I V E Q N Y G S T E G Y H G Q S G I L E E T E D T N P P G I I L G G S G N V E T H E D T R N P  
 H L M G I G G G L A G E S G E T T P K P G Q T G G Q G P V I E T T E D T Q K G M S G Q S G G T I E S E N T K K P E V M I G G  
 Q G Q T I E T T E D T Q K G M S G Q S G G T I E S E D T K K P E V M I G G Q Q I I D F S E N T Q S G M S G Q S G D T V I  
 E D T K K S E I I I G G Q Q I I D F S E D T Q P G M S G Q S G G T I V E D T K K P T P K P K P A P A P I V N D E K P N K  
 G T H L P Q T S D M K Q L T L S I I G A M S M L L V L C L S L F K R P S K K D

and SFS comprising an amino acid sequence as shown in SEQ. ID. NO: 3 below:

15 M R K T E G R F R T W K S K K Q W L F A G A V V T S L L G A A L V F G G L L G S L G G  
 S S H Q A R P K E Q P V S S I G D D D K S H K S S S D S M V S R P P K K D N L Q P K P S D Q P T N H Q H Q A T S P S  
 Q P T A K S S G H H G N Q P Q S L S V N S Q G N S S G Q A S E P Q A I P N Q G P S Q P L G L R G G N S S G S G H H H  
 20 Q P Q G K P Q H L D L G K D N S S P Q P Q P K P Q G N S P K L P E K G L N G E N Q K E P E Q G E A G P P L S G  
 L S G N N Q G R P S L P G L N G E N Q K E P E Q G E A G P P S T P N L E G N N R K N P L K G L D G E N K P K E  
 25 D L D G K G L S G E N D E S P K L K D E H P Y N H G R R D G Y R V G Y E D G Y G G K K H K G D Y P K R F D E S S P K  
 E Y N D Y S Q G Y N D N Y G N G Y L D G L A D R G G K R G Y G Y S Y N P D.

Suitably, the afore-mentioned antigenic compositions also comprise at least one further antigen that comprises at least part of an extracellular matrix-binding protein of *Streptococcus equi* and said at least part of said protein comprises an antigenic epitope or an 30 antigenic determinant of *Streptococcus equi* and wherein said protein is comprised of a

protein that is designated SEC and comprises an amino acid sequence as shown in SEQ. ID. NO: 4 below.

SEQ. ID. NO: 4:

1 27  
5 LKQLTKIVSVVLLLVTLSASLHKVRATNLSDNITSLTVA  
SSSLRDGERTTVKVAFDD  
KKQKIKAGDTIEVTWPTSGNYYIQGFNKTIPNIRGV  
DVGTLVTLDAVFTFNQNIE  
TMHDVSGWGEFDITVRNVTQTTAETSGTTVKVGN  
RTATITVTKPEAGTGTSSFYK  
TGDMQPNDTERVRWFLLINNNKEWVANTV  
TEDDIQGGQTLDMSSFDITVSGYRNE  
RFVGENALTEFHTTFPNSVITATDNHISV  
RLDQYDASQNTVNIAYKTKITDFDQKEFA  
10 NNSKIWYQILYKDQVSGQESNHQVANINANG  
GVDGSRYSFTVKKIWNDKENQDGK  
RPKTTIVQLYANDQKVNDKTI  
ELSDTNSWQASFGKLDKYDSQNQKITYSV  
KEVMVP  
VGYQSQVEGDSGVGFTITNTY  
TPEVISITGQKTWDDRENQDGKRP  
KEITVRL  
LANDA  
ATDKVATASEQTGWKY  
TFTNLPKY  
KDGKQITY  
TIQEDPVADY  
TTTIQGF  
DITNH  
HEV  
15 ALTS  
LKV  
IKV  
WNDK  
DDYYH  
KRP  
KEIT  
ILLK  
ADGK  
VIREH  
QMTPD  
QQGK  
WEY  
TFD  
QL  
PVY  
QTG  
KKI  
SIEEK  
QVAGY  
QAPV  
YEV  
DEGLK  
QVT  
VTN  
LPSY  
KLP  
DTGG  
QGV  
K  
WYLL  
IGGG  
FIIV  
AIL  
VLIS  
LYQ  
KHK  
RHN  
MSKP

657

The antigens of the present antigenic compositions may comprise the entire amino acid sequence of said protein or may comprise a fragment or analog thereof. Suitable fragments are N-terminal fragments of EAG and FNZ. An antigen derived from SFS may be comprised of a C-terminal part of SFS and an antigen derived from SEC may be comprised of a collagen-binding part of SEC.

A preferred antigenic composition of the present invention contains all the afore-said antigens EAG, FNZ, SFS, and SEC. Preferably, an N-terminal part of each of EAG and FNZ designated FNZN and EAG4B, respectively, a C-terminal part of SFS designated SFSC1 and a collagen-binding part of SEC designated SEC 2.16 are used in said composition.

From the above, it is evident that the present antigens that are derived from proteins of *Streptococcus equi* may comprise the entire protein, a fragment of said protein or an analog of said protein. Thus, the present invention is not limited to the fragments of proteins that are specifically disclosed herein.

A further embodiment of the present invention is concerned with a vaccine composition for protecting non-human mammals against infection of *Streptococcus equi*, which comprises an antigenic composition as disclosed above as immunizing component, and a pharmaceutically acceptable carrier.

35 Suitably, the present vaccine composition comprises an antigenic composition that contains all the afore-said antigens as immunizing component. Optionally, one or more of these antigens are comprised of analogs of said proteins or fragments thereof, e.g. N-terminal or C-terminal fragments.

The vaccine composition may comprise further components, such as an adjuvant. Suitably, said adjuvant stimulates systemic or mucosal immunity. Such adjuvants are well known in the art.

According to a suitable embodiment, the vaccine composition is a vaccine that 5 protects susceptible mammals, suitably horses, against strangles caused by *Streptococcus equi* subsp. *equi*.

The vaccine composition of the present invention is provided in a physiologically administrable form. Suitably, it is administrable by subcutaneous or intranasal inoculation. Suitably, the vaccine composition of the present invention stimulates serum, mucosal and/or 10 bronchial lavage antibody responses directed to *Streptococcus equi* antigens in mammals susceptible to *Streptococcus equi*, suitably horses.

The present invention is also related to a method for producing an antigen to be used in an antigenic composition of the present invention, which method comprises

- (a) providing a DNA fragment encoding said antigen and introducing said fragment 15 into an expression vector;
- (b) introducing said vector, which contains said DNA fragment, into a compatible host cell;
- (c) culturing said host cell provided in step (b) under conditions required for expression of the product encoded by said DNA fragment; and
- 20 (d) isolating the expressed product from the cultured host cell.

Preferably, said method further comprises a step (e) wherein the isolated product from step (d) is purified, e.g. by affinity chromatography or other chromatographic methods known in the art.

A further embodiment of the present invention is concerned with a method for 25 preparation of a vaccine of the present invention, which vaccine contains as immunizing component an antigenic composition as disclosed above, said method comprising mixing said antigenic composition and a pharmaceutically acceptable carrier.

The present invention is also related to a method for the production of an antiserum, said method comprising administering an antigenic preparation of the present invention to an 30 animal host to produce antibodies in said animal host and recovering antiserum containing said antibodies produced in said animal host.

Moreover, the present invention is concerned with a method of prophylactic or therapeutic treatment of *S. equi* infection in non-human mammals, suitably horses, comprising

administering to said mammal an immunologically effective amount of a vaccine or an antiserum of the present invention.

A suitable embodiment of the present invention is concerned with a method for protecting horses against *Streptococcus equi* infection, which comprises inoculating a horse 5 subcutaneously or intranasally or both subcutaneously and intranasally with a vaccine of the present invention to induce an immune response against *Streptococcus equi* in said horse. Suitably, an immune response in the form of IgG and IgA antibodies in the nasopharyngeal mucus is thereby induced in said horse.

The present invention describes the composition of a vaccine comprising one or 10 several antigen components which have been prepared according to the present method using *E. coli* as host cells. The source of these antigens might also be the native bacteria, if methods are developed for expression and purification thereof. Alternatively, the antigens of the present invention can also be produced according to methods that are based on fusion strategies where various parts of the respective antigen are recombined resulting in a fusion 15 protein consisting of parts from different antigens. This fusion strategy could also be suitable for introducing immune reactive part(s), e.g. T-cell epitopes or attenuated toxins (or parts thereof), thereby introducing other features suitable for optimizing the antigen presentation or localization.

## EXPERIMENTAL PART

### 20 Example 1. Preparation of antigens derived from EAG

A gene similar to the afore-mentioned *zag* gene (the GeneBank accession number for the nucleotide sequence of *zag* is U2582) from *S. equi* subsp. *zooepidemicus* but present in subsp. *equi* has been described in Lindmark et al (1999) (Ref. 11) and Lindmark (1999) (Ref. 13). This gene that is present in subsp. *equi* and encodes the afore-mentioned protein EAG is 25 hereafter called *eag*. To clone and express a part of this gene encoding an N-terminal fragment (EAG4B) of EAG in *E. coli*, the following work was done.

#### Construction

##### The primers

OZAG43B: 5'-TTT TCT CGA GCT ACG GTA GAG CTG ATA AAA TCT C-3'

30 (SEQ. ID. NO: 5) and

OZAG15: 5'-TCA GCC ATG GCT CTA GAT GCT ACA ACG GTG TT-3'

(SEQ. ID. NO: 6)

were used to PCR-amplify a DNA-fragment corresponding to amino acid residues 34-262 in protein EAG using *S. equi* subspecies *equi* 1866 chromosomal DNA as a template. The PCR-

product was digested with *Nco*I and *Xho*I and ligated into the pTYB4-vector obtained from New England Laboratories (NEB Inc) digested with the same enzymes. One  $\mu$ l of the ligation mixture was transformed into *E. coli* ER2566. Correct transformants were identified by colony screening using horse radish peroxidase-labeled human serum albumin. The clone 5 chosen for further work was sequenced from both directions to verify the correct insertion into the vector and the presence of a stop codon between EAG and the intein-affinity tag.

#### **HSA-column**

50 mg human serum albumin (HSA, Sigma) were immobilized on a 5 ml HiTrap NHS-activated column (Pharmacia Biotech) according to the manufacturer's instructions.

10 Based on the absorbance at 280 nm, 65% of the HSA was estimated to be immobilized on the column.

#### **Expression and purification – general**

16 ml of an over night culture of EAG4B in *E. coli* ER2566 were inoculated into 2 liters of LB-medium supplemented with 50  $\mu$ g ampicillin per ml. The culture was incubated at 15 37°C until an  $OD_{600}$  in the range of 0.5-0.6 was reached. Thereafter, IPTG was added to a final concentration of 0.5 mM and the culture was incubated at 25°C over night.

The culture was harvested and the pellet was washed once in 50 ml PBS-D before it was resuspended in 40 ml of 20 mM Na phosphate, pH 7.0 (binding buffer). The resuspension was divided into two tubes and 10 mg lysozyme were added to each tube. After 2 hrs. of 20 incubation at 37°C, the tubes were frozen at -20°C over night.

The lysate was sonicated to fragmentize the DNA, centrifuged and the supernatant was filtered through a 0.45  $\mu$ m filter.

The column was washed with 30 ml binding buffer before the sample (corresponding to 1 liter of the over night culture) was applied onto the column. Thereafter, the column was 25 washed with 50 ml binding buffer and the  $A_{280}$  value was confirmed to be below 0.010. Bound protein was eluted in 0.1 M glycine-HCl, pH 3.0. The first 4.5 ml were not collected while the next 10 ml were collected as 1 ml fractions, which were neutralized by addition of 100  $\mu$ l 1 M Tris-buffer, pH 8.0. Then, the column was regenerated by addition of binding buffer and stored at + 4°C in the presence of sodium azide.

30 The  $A_{280}$  value was determined and the fractions containing protein, normally fractions 1 to 5 or 6 were pooled and dialyzed 3 times against 4 liters of PBS-D. Thereafter, the protein was concentrated by extracting water from the dialysis tubing with PEG 30 000.

From an over night culture of 2 liters, this procedure yielded 5-6 mg of an N-terminal fraction of the EAG protein, designated EAG4B and comprising the amino acid sequence recited above as SEQ. ID. NO: 1. In this sequence, the first amino acid presented in bold originates from the vector.

5 The nucleotide sequence of the fragment encoding EAG4B where the coding sequence starts from nucleotide 38(A) and ends at nucleotide 577(C) is shown below (SEQ. ID. NO: 7).

SEQ. ID. NO: 7:

	10	20	30	40	50
10					
15	1 AAATAATTTGTTAACCTTAAGAAGGAGATATAACCATGGCTCTAGATG	51 CTACAAACGGTGTAGAGCCTACAACAGCCTTCATTAGAGAAGCTGTTAGG	101 GAAATCAATCAGCTGAGTGATGACTACGCTGACAATCAAGAGCTTCAGGC	151 TGTTCTTGCTAATGCTGGAGTTGAGGCACTTGCTGCAGATACTGTTGATC	201 AGGCTAAAGCAGCTTGACAAAGCAAAGGCAGCTGTTGCTGGTGTTCAG
20	251 CTTGATGAAGCAAGACGTGAGGCTTACAGAACAAATCAATGCCCTTAAGTGA	301 TCAGCACAAAAGCGATCAAAAGGTTCAGCTAGCTCTAGTTGCTGCAGCAG	351 CTAAGGTGGCAGATGCTGCTTCAGTTGATCAAGTGAATGCAGCCATTAAT	401 GATGCTCATACAGCTATTGCGGACATTACAGGAGCAGCCTTGGAGGC	451 TAAAGAAGCTGCTATCAATGAACCTAAAGCAGTATGGCATTAGTGATTACT
25	501 ATGTGACCTTAATCAACAAAGCCAAACTGTTGAAGGTGTCAATGCGCTT	551 AAGGCAAAGATTATCAGCTCTACCGTAGCTCGAGCCGGTGCTTG			

#### Example 2. Preparation of antigens derived from SFS

The SFS protein from *S. equi* subsp. *equi* has previously been described by Lindmark and Guss (1999) (Ref. 12) and in WO 00/37496. The GeneBank accession number for the nucleotide sequence of the *sfs* gene is AF 136451.

A C-terminal fragment of this protein was produced as follows:

The 3' end of the *sfs* gene was PCR amplified using the Taq DNA polymerase (Amersham) and chromosomal DNA from *S. equi* strain 1866 as template and the synthetic oligonucleotides

OSFS25: 5'-GGTCCCATGGCAACTCCGAATTAGAAGGA-3' (SEQ. ID. NO: 8) and OSFS23: 5'-CAGACTCGAGGTCGGGATTGTAAGAATAG-3' (SEQ. ID. NO: 9) as primers. The PCR procedure was performed in 100 µl under standard conditions as regards buffer, template and primer concentration. The PCR cycles were performed under the following conditions: 1 min. of denaturation at 94°C, 30 sec. of annealing at 40°C, and 2 min. of extension at 72°C, which were repeated as 25 cycles. After amplification, the PCR product

was purified using phenol extractions, chloroform extractions and EtOH precipitation. The purified DNA was cleaved with the restriction enzymes *Nco*I and *Xho*I after which the DNA was purified as described above.

The DNA obtained above was ligated into the plasmid vector pTYB4 [(New England Biolabs, Beverly, MA, USA (NEB Inc.)), which previously had been digested with the same restriction enzymes and treated with alkaline phosphatase. After ligation (using the ReadyToGo ligation kit, Amersham), the DNA sample was electrotransformed into the *E. coli* strain ER2566 and spread on LAA-plates (Luria Bertani agar plates supplemented with ampicillin, final conc. 50 µg/ml). After incubation over night at 37°C, clones harboring plasmids with inserts were isolated and the presence of the correct insert was verified by DNA sequencing. One of the clones thereby obtained, called SFSC1, was chosen for production of the C-terminal part of SFS.

The vector used is a part of an *E. coli* expression and purification system called IMPACT<sup>T</sup> T7 (NEB Inc.). Briefly, following the manufacturer's instructions, the clone SFSC1 was grown at 37 °C in Luria Bertani growth medium supplemented with ampicillin (final conc. 50 µg/ml). At an optical density (OD<sub>600</sub>) of ~ 0.6, the growth medium was supplemented with IPTG (final conc. 0.4 mM) and the growth temperature was shifted to 20°C. After incubation over night, the cells were harvested and resuspended in a buffer [20 mM Tris-HCl (pH 8.0), 500 mM NaCl, 0.1 mM EDTA, and 0.1 % Triton X100], lysed by freezing and thawing, and after centrifugation, the supernatant was sterile filtered and applied to a chitin column.

The column was extensively washed using the same buffer as above and was subsequently treated with cleavage buffer [20 mM Tris-HCl (pH 8.0), 50 mM NaCl, 0.1 mM EDTA, and 30 mM dithiothreitol (DTT)]. The reducing conditions in the cleavage buffer induce an intein-mediated self-cleavage that releases the SFS part from the column while the intein-chitin-binding part remains bound. The eluted sample containing the product SFSC1 was dialysed against phosphate-buffered saline [PBS; 137 mM NaCl, 2.7 mM KCl, 10 mM Na<sub>2</sub>HPO<sub>4</sub>, 1.4 mM KH<sub>2</sub>PO<sub>4</sub> (pH 7.4)] and concentrated. The amount of protein obtained was determined and the quality checked using SDS-PAGE.

After the intein part has been cleaved off from SFSC1, the purified protein has an amino acid sequence as recited below as SEQ. ID. NO: 10 except that the amino acid residues in bold are residues that correspond to the amino acid residues encoded by the pTYB4 vector, while remaining amino acid residues originate from the SFS protein.

SEQ. ID. NO: 10:

MATPNLEGNN RKNPLKGLDG ENKPKEDLDG KGLSGENDES PKLKDEHPYN  
HGRRDGYRVG YEDGYGGKKH KGDYPKRFDE SSPKEYNDYS QGYNDNYGNG  
YLDGLADRGG KRGYGYSYNP DLEPG.

**Example 3. Preparation of antigens derived from FNZ**

5 Protein FNZ from *S. equi* subsp. *zooepidemicus* has previously been described by Lindmark et al (1996) (Ref. 9), Lindmark (1999) (Ref. 13), Lindmark et al (1999) (Ref. 11) and Lindmark et al (2001) (Ref. 14). The EMBL accession number for the nucleotide sequences of the *fnz* gene is X99995. A similar truncated protein called FNE is also expressed by strains of *S. equi* subsp. *equi* [Lindmark et al (2001), Ref. 14]. The GeneBank accession 10 number for the nucleotide sequence of the *fne* gene is AF360373. The construction of a clone called pT2fnzN encoding the N-terminal part of FNZ has previously been described by Lindmark et al 2001 (Ref. 14). Briefly, the clone pT2fnzN was constructed as follows using PCR amplification with the forward primer  
OFNZ1: 5'-ACCATGGCTAGCGCAGAGCAGCTTATTATGGGT-3' (SEQ. ID. NO: 11),  
15 and the reverse primer  
OFNZ2: 5'- ATACCCGGGATATCCTTCGGTACTACCATAGT-3' (SEQ. ID. NO: 12). Chromosomal DNA from subsp. *zooepidemicus* strain ZV was used as the template and the 5' end of the *fnz* gene was amplified. The PCR fragment obtained was cleaved with restriction endonucleases *Nhe*I and *Sma*I, followed by ligation into the corresponding restriction 20 endonuclease sites in the expression vector pTYB2 (NEB). The ligated DNA was electrotransformed into *E. coli* strain ER2566. Plasmids harboring inserts were isolated from transformants and verified by DNA sequencing. One such clone is designated pT2fnzN. The production in *E. coli* and purification of the N-terminal part of FNZ, called FNZN, were performed as described above for protein SFSC1 and these steps are also described in  
25 Lindmark et al (2001) (Ref. 14).

The amino acid sequence of FNZN comprises the amino acid residues 32-337 of the amino acid sequence of the protein FNZ recited above as SEQ. ID. NO: 2. The corresponding amino acid sequence of FNZN is shown below as SEQ. ID. NO: 13.

30 SEQ. ID. NO: 13

M A S A E Q L Y Y G W N D G T R Q S S P  
Y F L Y V S P K N A P K R E L K D E Y V  
V Y C F N K K L Y W P D Q W E S I Y S N  
F N D I R S P Y N D L P V Y E K K L G Y  
35 D G I F K Q Y A P D Y K K D I S D I A S

	A	L	V	A	V	L	S	N	G	Y	P	T	N	K	S	Q	L	S	T	S
	Y	H	L	N	N	D	S	S	R	K	V	T	Q	L	A	I	W	Y	F	S
	D	S	L	T	K	E	Y	L	K	D	T	G	G	Y	N	L	N	D	M	E
	K	K	A	L	D	F	L	I	S	K	G	E	D	S	K	L	K	S	E	Q
5	S	N	Y	S	L	D	I	Y	V	Y	Q	S	G	G	H	D	H	M	K	D
	Y	Q	N	L	L	G	S	T	L	I	P	K	E	P	L	K	P	Q	L	G
	G	F	S	G	H	N	G	N	G	L	S	G	L	E	G	G	S	S	G	S
	Q	E	T	N	E	D	G	K	K	G	L	I	G	F	H	G	G	L	S	G
10	S	E	G	K	R	D	P	L	P	G	L	K	G	E	A	G	A	P	D	T
	P	Q	K	P	N	D	P	L	Q	G	L	E	G	G	N	S	P	I	V	E
	Q	N	Y	G	S	T	E	G	Y	G										

In the previously recited amino acid sequence (SEQ. ID. NO: 2) of protein FNZ, amino acids shown in bold originate from the protein FNZ in *S. equi* subspecies *zooepidemicus*, while those shown in normal type are derived from the expression vector construct. In SEQ. ID. NO:13, the first three and the last amino acid residues originate from the vector.

**Example 4. Preparation of antigens derived from the protein SEC**

The genome of *S. equi* is accessible ([www.sanger.ac.uk](http://www.sanger.ac.uk)) for computer analysis. By using earlier published sequences of virulence factors or potential virulence factors from pathogenic streptococci and staphylococci, it is possible to screen the genome of *S. equi* for the presence of similar genes. By using the soft ware program BLAST and searching for open reading frames encoding a protein similar to the collagen-binding protein CNA of *Staphylococcus aureus* (WO 92/07002), it was possible to identify a hypothetical gene of 1971 nucleotides in the *S. equi* genome encoding a protein of 657 amino acids. (Ref. 25, Lannergård et al. (2003)). This protein, hereafter termed protein SEC (SEQ. ID. NO: 4), encodes an N-terminal signal sequence (amino acid sequence 1-26) followed by a region, which displays collagen-binding. The C-terminal part of protein SEC contains all typical features of cell surface proteins from streptococci, like putative wall anchoring and membrane spanning regions, as well as a motif corresponding to the LPDTG motif (SEQ. ID. NO: 14) of the protein SEC.

The nucleotide sequence of the *sec* gene (later renamed to *cne*) encoding the protein SEC (the accession no. of the *cne* is AY193773). is shown below as SEQ. ID. NO: 15.

SEQ. ID. NO: 15:

1

TTGAAACAACTGACAAAGATCGTTAGTGTGGCTTGTGCTGGCTTACCCCTAG  
 TGCTAGCCTGCACAAGGTTGGGCAACTAATCTTAGTGACAACATCACATCATTG  
 ACGGTTGCTCTTCATCACTCCGAGATGGAGAGAGAACGACGGTAAAGGGTGCCTG  
 TTGATGACAAAAAAACAGAAAATCAAGGCAGGGGATACGATAGAGGTACACCTGGC  
 CTACAAGTGGTAATGTCTACATTCAAGGGCTTAATAAAACCATAACCGCTTAATAT  
 TAGAGGGGTAGATGTTGGTACCTGGAGGTACGCTAGACAAGGCTGTTTCACA  
 TTCAATCAAATATTGAAACAAATGCATGATGTCCTGGTTGGGAGAGGTTGATA  
 TTACTGTTAGAAATGTGACACAAACCACCGCTGAAACATCAGGAACGACCAACAG  
 TAAAGGTAGGCAATCGCACTGCTACTATCACTGTTACTAAGCCTGAGGCAGGCAC  
 TGGTACCAAGCTCATTATTATAAGACTGGTATATGCAGCCCAATGATACTGAG  
 CGTGTGAGATGGTTCCTGCTGATTAACAACAAACAAGGAATGGTGGCCAATACTG  
 TTACAGTCGAAGACGATAATTCAAGGTGGTCAAACCTTGGATATGAGCAGCTTGA  
 CATCACCGTATCTGGTTATCGTAACGAGCGCTTCGTTGGGAAAACGCTCTGACA  
 GAGTTTCATACAACATTTCAAATTCTGTCATTACGGCAACAGATAATCACATTA  
 GTGTGCGGTTAGATCAATATGATGCTCACAACAAACTGTCACACATTGCTTATAA  
 GACAAAGATAACGGACTTGACCAAAAAAGAATTGCCAACAAACAGTAAAATCTG  
 GTACCAAGATTTATACAAGGATCAGGTATCGGGTCAAGAGTCAAACACCACCAAGTA  
 GCCAATATCAATGCTAACGGCGGGTTGATGGCAGTCGCTATACCAGCTTACTG  
 TCAAGAAAATTGGAATGACAAGGAAAATCAAGACGGTAAGCGTCCAAAGACTA  
 TTACTGTTAGCTTACGCCAATGATCAGAAAGTTAATGATAAGACCATTGAATT  
 GAGTGTACTAAATAGCTGGCAAGCAAGTTGGTAAGCTGGATAAGTATGACAGT  
 CAGAACCAAAAAATTACCTACAGTGTCAAGGAAGTGTAGGGTCTGTTGGCTACC  
 AATCGCAGGTTGAGGGGATAGTGGAGTAGGATTACCAACACCTATAC  
 ACCAGAGGTATTACGATTACCGGTCAAAAAACTGGGACGACAGGGAAAACCA  
 AGACGGTAAACGTCCTAACGGAGATTACGGTTGTTATTGGCAAATGACGCTGCA  
 ACTGACAAGGTAGCAACTGCTTCAGAGCAAACCGGCTGGAAGTATACATTACCA  
 ATCTACCGAAATACAAAGATGGTAAACAGATCACCTACACGATCCAAGAGGACC  
 CTGTGGCAGATTACACCACAACCATTCAAGGGATTGATATTACCAATCATCATGA  
 GGTAGCCTTGACCAGCCTAACGGTCAAGGTTGGATAGATAAGGACGATTAT  
 TACCAAAACGTCCAAGGAGATTACCAATTGCTAAAGGCAGATGGCAAGGTGA  
 TTCGTGAACATCAGATGACACCCGGATCAGCAAGGAAAATGGAATACACCTTG  
 ACCAGCTGCCGGTCTATCAGACAGGCAAGAAAATCAGCTACAGCATTGAGGAAA  
 AACAGGTTGCTGGCTATCAAGCCCTGCTATGAGGTGATGAAGGCTTGAAGCA  
 GGTCACTGTAACCAACACCCTAACCCAAAGCTACAAGCTGCCTGACACCGGAGGA  
 CAAGGAGTGAATGGTACCTGTTAACCGTGGCGGTTATCATCGTCGCAATCC  
 TTGTACTGATCAGCCTTATCAAAAACACAAGGCCATAACATGTCAAAACCA

1971

5

### Construction of recombinant protein SEC expressing collagen-binding activity

To express and purify the collagen-binding activity of protein SEC, two different constructions were made. The first construct, encoding the major part of protein SEC was  
 10 made as follows. The primers

OSEC1:5: 5'-CATGCCATGGCAACTAATCTTAGTGACAACAT-3' (SEQ. ID. NO: 16)  
and

OSEC3:3: 5'-CCGCTCGAGCTTGTAGCTGGGTTAAGGGTGT -3' (SEQ. ID. NO: 17)  
were used to PCR-amplify a DNA-fragment corresponding to a sequence from amino acid no.

27 to amino acid no. 615 in protein SEC (SEQ. ID. NO: 4) using *S. equi* subspecies *equi* 1866 chromosomal DNA as a template.

The second construct encoding the N-terminal part of protein SEC was made as follows. The primers OSEC1:5 and

5 OSEC2:3: 5'-CCGCTCGAGAAAGCTGGTATAGCGACTGCCAT-3' (SEQ. ID. NO: 18) were used to PCR-amplify a DNA-fragment corresponding to a sequence from amino acid no. 27 to amino acid no. 328 in protein SEC (SEQ. ID. NO: 4) using *S. equi* subspecies *equi* 1866 chromosomal DNA as a template.

Both PCR amplifications were performed using ReadyToGo<sup>TM</sup> PCR beads

10 (Amersham Pharmacia Biotech Inc) and the PCR apparatus MiniCycler<sup>TM</sup> (MJ Research, Inc, MA, USA) using a program comprising: step 1: 95°C, 1 min; step 2: 95°C, 30 sec.; step 3: 46°C, 15 sec.; and step 4: 72°C, 2 min; repeated as 25 cycles from step 2 to step 4. After PCR amplification, the respective PCR-products were purified and digested with restriction endonucleases *Nco*I and *Xho*I and ligated into the pTYB4-vector (NEB) digested previously 15 with the same enzymes. One µl of the respective ligation mixture was transformed into *E. coli* ER2566. After IPTG induction, *E. coli* clones expressing collagen-binding were identified by colony screening using <sup>125</sup>I-labeled collagen (collagen S, type I obtained from Boeringher Mannheim). Several clones of both types of constructions expressing collagen-binding were identified and further characterized. One of these clones called pSEC 2.16 harboring the PCR- 20 fragment and originating from the PCR using OSEC 1:5 and 3:3 was chosen after DNA sequencing of the insert for production of a recombinant collagen-binding protein called protein SEC 2.16 and comprising amino acid residues 27 - 616 of SEQ. ID. NO: 4. The nucleotide sequence of this insert encoding the recombinant collagen-binding protein SEC2.16 is shown below as SEQ. ID. NO: 19. The nucleotides shown in bold represent 25 nucleotides originating from the vector.

SEQ. ID. NO: 19: Insert of pSEC2.16

ATGGCAACTAATCTTAGTGACAACATCACATCATTGACGGTTGCTTCTTCATCACTCCGAGATGGA  
GAGAGAACGACGGTAAAGGGTGCCTTGTATGACAAAAAAACAGAAAATCAAGGCAGGGGATACGA  
TAGAGGTACCTGGCCTACAAGTGGTAATGTCTACATTCAAGGGCTTAATAAAACCATAACGCTTA  
ATATTAGAGGGTAGATGTTGGTACCTTGGAGGTACCGCTAGACAAGGCTGTTTCACATTCAAT  
CAAAATATTGAAACAAATGCATGATGTCTCTGGTGGGGAGAGTTGATATTACTGTTAGAAATGTG  
ACACAAACCACCGCTGAAACATCAGGAACGACCACAGTAAAGGTAGGCAATCGCACTGCTACTAT  
CACTGTTACTAAGCCTGAGGCAGGCAGTGGTACCAAGCTCATTATTATAAGACTGGTGTATTCA  
GCCCAATGATACTGAGCGTGTGAGATGGTCTGCTGATTAACAACAACAAGGAATGGGTGGCC  
AATACTGTTACAGTCGAAGACGATATTCAAGGTGGTAAACCTTGGATATGAGCAGCTTGACATC  
ACCGTATCTGGTTATCGTAACGAGCGCTTCGTTGGGGAAAACGCTCTGACAGAGTTCATACAAC  
ATTCCAATCTGTCATTACGGCAACAGATAATCACATTAGTGTGCGGGTAGATCAATATGATGC  
CTCACAAAACACTGTCAACATTGCTTATAAGACAAAGATAACGGACTTGA  
CCAAAAAGAATTG

CAACAACAGTAAAATCTGGTACCAGATTTATACAAGGATCAGGTATCGGGTCAAGAGTCAAACC  
 ACCAAGTAGCCAATATCAATGCTAACGGCGGGGTTGATGGCAGTCGCTATACCAAGCTTACTGTC  
 AAGAAAATTGGAATGACAAGGAAAATCAAGACGGTAAGCGTCAAAGACTATTACTGTTAGCCT  
 TACGCCAATGATCAGAAAGTTAATGATAAGACCATTGAATTGAGTGATACTAATAGCTGGCAAGCA  
 AGTTTGGTAAGCTGGATAAGTATGACAGTCAGAACCAAAAAATTACCTACAGTGTCAAGGAAGT  
 GATGGTTCTGTTGGCTACCAATCGCAGGTTGAGGGGGATAGTGGAGTAGGATTACCATTACCA  
 ACACCTATACACCAGAGGTCAATTAGCATTACCGTCAAAAAACTTGGGACGACAGGGAAAACCAA  
 GACGGTAAACGTCTAACGGAGATTACGGTCTTATTGGCAAATGACGCTGCAACTGACAAGGT  
 AGCAACTGCTTCAGAGCAAACCGGCTGGAAGTATACATTACCAATCTACCGAAATACAAAGATG  
 GTAAACAGATCACCTACACGATCCAAGAGGACCCCTGTCAGATTACACCACAACCATTAGGGA  
 TTTGATATTACCAATCATCATGAGGTAGCCTGACCAGCCTAAAGGTCAAGGTTGGAATGAT  
 AAGGACGATTATTACCATAAACGTCCAAGGAGATTACCATTTGCTAAAGGCAGATGGCAAGGTG  
 ATTCGTGAACATCAGATGACACCGGATCAGCAAGGAAAATGGAATACACCTTGACCAGCTGC  
 CGGTCTATCAGGCAGGCAAGAAAATCAGCTACAGCATTGAGGAAAACAGGTTGCTGGCTATCAA  
 GCCCCTGTCTATGAGGTTGATGAAGGCTGAAGCAGGTCACTGTAACCAACACCCCTAACCAAG  
 CTACAAGCTCGAGCCCCGGG

The amino acid sequence of the recombinant collagen-binding protein SEC2.16  
 encoded by the insert of pSEC2.16 is shown below as SEQ. ID. NO: 20. The amino acids  
 shown in bold represent amino acids originating from the vector.

5

SEQ. ID. NO: 20: Protein SEC2.16

MATNLSDNITSLTVAASSLRDGERTVKVA**F**DDKKQ**K**IKAGDTIEVTWPTSGNVYIQGFNK**T**IPLNIRGV  
 DVGTLEVTLKAVFTFNQNIETMHDVSGWGEFDITVRNVTQTTAETSGTTVKG**N**R**T**ATITVTKPEA  
 GTGTSSFYYKTGDIQPNDTERVRWFLLINNNKEWVANTVTVEDDIQGGQ**T**LD**M**SSFDITVSGYRNER  
 FVGENALTEFHTTFPNSVITATDNHISVRLDQYDASQNTVNIA**Y**TK**K**ITDFD**Q**KEFANNS**K**I**W**Y**Q**IL**Y**KD  
 QVSGQESNHQVANINANGGV**D**GSRYTSFTVK**K**I**W**ND**K**ENQDG**K**R**P**K**T**ITV**Q**LY**Y**ND**Q**KVND**K**TI**E**LS  
 DTNSWQASF**G**KLD**K**Y**D**SQ**N**Q**K**ITY**S**KE**V**M**V**P**G**Y**Q**S**Q**VE**G**DS**G**VG**F**ITNTY**T**PE**V**ISIT**G**Q**K**T**W**D  
 DRENQDG**K**R**P**KE**I**TV**R**LL**A**AT**D**K**V**AT**A**SE**Q**T**G**W**K**Y**T**FT**N**LP**K**Y**K**D**G**K**Q**ITY**T**IQ**E**DP**V**ADY**T**TI**Q**  
 GF**D**IT**N**H**E**VAL**T**SL**K**VI**K**V**W**ND**K**DDYY**H**K**R**PE**I**LL**K**AD**G**K**V**IRE**H**Q**M**TPD**Q**Q**G**K**W**ET**F**D**Q**LP  
 VY**Q**AG**K**K**I**SY**S**IE**E**K**Q**V**A**GY**Q**AP**V**Y**E**V**D**EG**L**K**Q**VT**V**T**N**LP**S**Y**K**LE**P**G

10 The other clone obtained above that was chosen for further studies was called pSEC 1.18 and  
 harbored the PCR-fragment originating from the PCR using OSEC 1:5 (SEQ. ID. NO. 16)  
 and 2:3 (SEQ. ID. NO. 18). After DNA sequencing of the insert of pSEC 1.18, this clone was  
 used for production of a recombinant collagen-binding protein called protein SEC 1.18. This  
 protein SEC 1.18 comprises the amino acid sequence shown below as SEQ. ID. NO. 22. The  
 15 corresponding nucleotide sequence is shown below as SEQ. ID. NO. 21. In this nucleotide  
 sequence, the nucleotides shown in bold represent nucleotides originating from the vector.

SEQ. ID. NO. 21: Insert of pSEC1.18

ATGGCAACTAATCTTAGTGACAACATCACATCATTGACGGTTGCTTCTTCATCACTCCGAGATGGA  
 GAGAGAACGACGGTAAAGGTTGCCTTGTAGACAAAAAAACAGAAAATCAAGGCAGGGGATACGA  
 TAGAGGTACCTGGCCTACAAGTGGTAATGTCTACATTCAAGGGCTTAATAAAACCATAACGCTTA  
 ATATTAGAGGGTAGATGTTGGTACCTGGAGGTACGCTAGACAAGGCTTTACATTCAAT  
 CAAAATATTGAAACAATGCATGATGTCCTGGTTGGGGAGAGTTGATATTACTGTTAGAAATGTG

ACACAAACCACCGCTGAAACATCAGGAACGACCACAGTAAAGGTAGGCAATCGCACTGCTACTAT  
CACTGTTACTAAGCCTGAGGCAGGCAGTGGTACCAAGCTCATTATTATAAGACTGGTATATGC  
AGCCCAATGATACTGAGCGTGTGAGATGGTCTGCTGATTAACAACAAGGAATGGGTGGC  
CAATACTGTTACAGTCGAAGACGATATTCAAGGTGGTCAAACCTGGATATGAGCAGCTTGACAT  
CACCGTATCTGGTTATCGTAACGAGCGCTTCGTTGGGAAAACGCTCTGACAGAGTTCATACAA  
CATTCCAATTCTGTCATTACGGCAACAGATAATCACATTAGTGTGCGGTTAGATCAATATGATG  
CCTCACAAAACACTGTCAACATTGCTTATAAGACAAAGATAACGGACTTGACCAAAAAGAATTG  
CCAACAACAGTAAAATCTGGTACCAAGATTTATACAAGGATCAGGTATCGGGTCAAGAGTCAAAC  
CACCAAGTAGCCAATATCAATGCTAACGGCGGGTTGATGGCAGTCGCTATACCAGCTTCTCGA  
GCCCGGG

The amino acid sequence of the recombinant collagen-binding protein SEC1.18 encoded by the insert of pSEC1.18 is shown below as SEQ. ID. NO: 22. The amino acids shown in bold represent amino acids originating from the vector.

5

SEQ. ID. NO: 22: Protein SEC1.18.

MATNLSDNITSLTVA~~SSSL~~RDGERTVKVA~~FDDKKQ~~KIKAGDTIEVTWPTSGNVYIQGFNKT~~IPLNIRGV~~  
DVG~~TLEV~~TLDKAVFTFNQNIETMHDVSGWGEFDITVRNVTQTTAETSGTTVKG~~NRT~~TATITVTKPEA  
GTG~~TSS~~FYYKTGDMQPNDTERVRWFLLINNNKEWVANTV~~T~~VEDDIQGGQ~~T~~LDMSSFDITVSGYRNER  
FVGENALTEFHTTFPNSVITATDNHISVRLDQYDASQNTVNIAYKTKITDFDQ~~K~~E~~F~~ANNSKIWYQILYKD  
QVSGQESNHQVANINANGV~~DGS~~RYTSFLEPG

#### Production of protein SEC 1.18 and protein SEC 2.16

10 Since the vector used to construct the pSEC1.18 and pSEC 2.16 is a part of the IMPACT-system (NEB), the expression and purification of protein SEC1.18 and protein SEC2.16 were performed as described for protein SFSC1 and protein FNZN. After purification, protein SEC 1.18 and protein SEC 2.16 were analysed by SDS-PAGE using the Phast-System (Amersham Pharmacia) and 8-25% gradient gels under reducing conditions  
15 (SDS buffer strips, boiling the samples in a sample buffer containing SDS and  $\beta$ -mercapto-ethanol before applying the samples on the gels). The results show that the purified protein SEC 1.18 and protein SEC 2.16, approx. had relative molecular weights corresponding to their calculated molecular weights.

#### Collagen-binding activity.

20 The purified proteins SEC 1.18 and SEC 1.16 were also analysed for their ability to bind to collagen. First, the proteins were run on an SDS-PAGE 8-25% gel (Amersham Pharmacia) under reducing conditions. After the electrophoresis was completed, the proteins were transferred (by diffusion) to a nitro-cellulose membrane. The membrane was blocked in a solution (PBS-T) containing PBS supplemented with TWEEN 20 (final concentration 0.5%)  
25 and casein 0.5% (final conc.) for 1 hr. at room temperature (RT). After washing with PBS-T (casein omitted), the membrane was transferred to a solution of PBS-T containing  $^{125}$ I-labeled

collagen. After 4 hrs. of incubation at RT under gentle agitation, the membrane was extensively washed using PBS-T (casein omitted) and subjected to autoradiography using Biomax MS (Kodak) film. The results showed that in contrast to unrelated proteins used as control (size marker proteins), both protein SEC 1.18 and protein SEC 2.16 bound collagen.

5       **Occurrence of the *sec* gene in *S. equi***

Using chromosomal DNA from various strains of *S. equi* subspecies *equi* and subspecies *zooepidemicus* as template and the primer combinations OSEC1:5 (SEQ. ID. NO: 16) and OSEC 2:3 (SEQ. ID. NO: 18) or OSEC1:5 (SEQ. ID. NO: 16) and OSEC 3:3 (SEQ. ID. NO: 17), the presence of the *sec* gene was investigated. The PCR conditions applied were

10      the same as described above in the section "Construction of recombinant protein SEC expressing collagen-binding activity". The result showed that the *sec* gene was present in all tested strains in both subspecies. This is an important finding, since a vaccine containing an antigen, which is present in both subspecies, gives the possibility to broaden the application of the invention resulting in a vaccine active against infections of both subspecies.

15

**Further search for vaccine components.**

The genome of *S. equi* is accessible ([www.sanger.ac.uk](http://www.sanger.ac.uk)) for computer analysis. As for protein SEC, it is possible to further screen the genome of *S. equi* for genes encoding earlier published sequences of virulence factors or potential virulence factors from pathogenic

20      streptococci and staphylococci. Use of the soft ware program BLAST resulted in the identification in *S. equi* of additional genes encoding potential virulence factors similar to *Streptococcus pyogenes*: C5a peptidase (gi:14195215, sp, P58099,SCA2 STRPY), SclB (emb: CAC33776.1), collagen-like protein similar to (ref: NP358996.1), speK (ref: NP 438166.1), exotoxin I (gi:16923748, gb:AAL31571.1, AF438524.1), streptodornase (emb:CAA59264.1),  
25      a DNA entry nuclease (ref: NP 346391.1), trypsin resistant surface T6 protein (sp:P18481, TEE6 STRPY), M and M-like proteins, other fibronectin-binding proteins and other.

Furthermore, the computer analyses revealed several hypothetical proteins displaying a structure which is found in cell wall-associated proteins in Gram-positive cocci (Schneewind et al 1995, Ref. 15). At the N-terminal end of these proteins, a signal sequence is found and at the C-terminal end, a motif LPXTG corresponding to the LPDTG motif (SEQ. ID. NO: 14) of the SEC protein is present, which is followed by a stretch of hydrophobic amino acids ending with a few charged amino acids constituting the membrane-spanning region. These proteins should also be considered as potential targets to be included in an efficient vaccine against *S. equi* infections.

Based on considerations as discussed above, two proteins have been prepared and evaluated as disclosed in the following Examples 5 and 6.

**Example 5. Preparation of protein SclC and protein SCLC1**

The amino acid sequence of *Streptococcus pyogenes* protein SclB was used to screen the genome of *S. equi* ([www.sanger.ac.uk](http://www.sanger.ac.uk)) using the software program BLAST. As a result of this screening, an open reading frame was identified, which encodes a protein called SclC, which is similar to SclB. The protein SclC comprises an amino acid sequence as shown as SEQ. ID. NO: 23 below.

SEQ. ID. NO: 23:

1

20 M T N K T K R T G L V R K Y G A C S A A I A L A A L A S L G A G K A V K A D  
Q P A A L K Y P E P R D Y F L H T R E G D V I Y D E D I K R Y F E D L E A Y  
L T A R L G G I D K K V E E A A Q K P G I P G P T G P Q G P K G D K G D P G  
A P G E R G P A G P K G D T G E A G P R G E Q G P A G Q A G E R G P K G D  
P G A P G P K G E K G D T G A V G P K G E K G D T G A T G P K G D K G E R  
25 G E K G E Q G Q R G E K G E Q G Q R G E K G E Q K P K G D Q G K D T K P  
S A P K A P E K A P A P K A P K A S E Q S S N P K A P A P K S A P S K S A A  
P T G Q K A A L P A T G E I N H P F F T L A A L S V I A S V G V L T L K G K  
K D

302

30 The nucleotide sequence of the gene sclC encoding protein SclC is shown below as SEQ. ID. NO: 24.

SEQ. ID. NO: 24:

35 1  
atgacaaacaaaacaaagcgtacaggattggtagcgaagtacggtgctgcctcagcagctatgccttagcagcttgcagccctgg  
gagcaggtaaaggcagttaaaggcagaccagccagcagcactaaaatatccagaaccttagagactatttccttcatactcgtaagggtat  
gttattttatgaggatataaaagatatttgaggatttagaaggctattaaacagcttagacttggggattgataaaaaagtagaagaa  
40 gctgccaaaagccaggtaatccaggctactggccctcaaggcttaagggagacaaggagatccagggtgcccctggtagcgc  
ggtccaggctggaccaaggcgatcggcgaagccggccaaggaggtgagcaaggcccagccggacaagctggagaacgtgg  
acccaaaaggagatccaggctccaggcttaagggtgatactggcagttggcttaagggtgatacc

ggagcaaccggaccaaaggagacaagggcaacgcggtaaaaaggcgagcaaggccaaactggcgaaaaaggcgagcaa  
ggccaaacgcggtaaaaaggcgagcaaaaaccaagggtatcaaggaaaagataaaaaaccatcagtcaccaagcacctgaaa  
aggctcctgcaccaaaggctcagagcagtcatctaattcataaggcaccagctcctaagtgcagcacaagcaaattcagc  
ggcaccaacaggtaaaaagcagccattaccagcaacaggaaatcaaccaccatttcacccttgagctcttagtgtcatcgcta  
5 gcgtaggcgtctaactctaaaagaaaaagactaa

909

To express and purify the major part of the SclC protein, the following construction was made. The primers

OSCL2:5: 5'- CATGCCATGGACCAGCCAGCAGCACTAAAATAT-3' (SEQ. ID. NO: 25)  
10 and OSCL3:3: 5'- CCGCTCGAGGGCTGCTTTGACCTGTTGGT-3' (SEQ. ID. NO: 26) were used to PCR-amplify a DNA-fragment corresponding to amino acid 38 to amino acid 269 in protein SclC using *S. equi* subspecies *equi* 1866 chromosomal DNA as a template. PCR amplification was performed using ReadyToGo<sup>TM</sup> PCR beads (Amersham Pharmacia Biotech Inc) and the PCR apparatus MiniCycler<sup>TM</sup> (MJ Research, Inc, MA, USA) using a  
15 program of step 1: 95°C, 1 min; step 2: 95°C 30 sec.; step 3: 50°C 15 sec.; step 4: 72°C, 1 min; repeated as 24 cycles from step 2 to step 4. After PCR amplification, the respective PCR-products were purified and digested with restriction endonucleases *Nco*I and *Xho*I and ligated into the pTYB4-vector (NEB) previously digested with the same enzymes. One µl of the respective ligation mixture was transformed into *E. coli* ER2566. *E. coli* clones harboring  
20 the *sclc* fragment were identified by DNA sequencing of the inserted fragments.

Of these clones, one clone called pSclC1 and harboring the PCR-fragment originating from the PCR using OSCL2:5 (SEQ. ID. NO: 25) and OSCL3:3 (SEQ. ID. NO: 26), was chosen for production of a recombinant SclC protein called protein SCLC1. This protein SCLC1 comprises the amino acid sequence shown below as SEQ. ID. NO: 27.

25

SEQ. ID. NO: 27:

**M D Q P A A L K Y P E P R D Y F L H T R E G D V I Y D E D I K R Y F E D L E**  
AYLTARLGGIDKKVEEAQKPGIPGPTGPQGPKGDKGDPGAPGERGPAGPKGDGTGEAGPRGEQGPGAGQAGERGPK  
30 **G D P G A P G P K G E K G D T G A V G P K G E K G D T G A T G P K G D K G**  
ERGEKGEQGQRGEKGEQGQRGEKGEQKPKGDQGKDTKPSAPKAPKAPKASEQSSNPKAPAPKSAPS  
35 **K A A L E P G**

35

In the peptide sequence of the insert of pSclC1 encoding the recombinant protein SCLC1 (SEQ. ID. NO: 27), the amino acids shown in bold represent amino acids originating from the vector.

#### Production of protein SCLC1

Since the vector used to construct the pSclC1 is a part of the IMPACT-system (NEB) the expression and purification of protein SCLC1 was performed as described for the other IMPACT-constructions (proteins SFSC1, FNZN, SEC 1.18 and SEC 2.16). After purification protein SCLC1 was analysed by SDS-PAGE using the Phast-System (Amersham Pharmacia) and 8-25% gradient gels under reducing conditions (SDS buffer strips, boiling the samples in a sample buffer containing SDS and beta-mercaptoethanol before applying the samples on the gels). The results showed that the purified protein had a relative molecular weight corresponding to its calculated molecular weight. The purified protein SCLC1 was then used to immunize mice and rabbits and was also used to screen convalescence sera from horses for 10 measurements of antibody titers against SCLC1. This is illustrated in Example 6 below.

**Example 6. Immunization of rabbits using protein SCLC1 and Western blot**

Protein SCLC1 was used to immunize a rabbit. This work was done at the company Agrisera (Vännäs, Sweden). Using an ELISA format, where the SCLC1 protein was immobilized in microtiter wells, the immune serum obtained was analysed for presence of 15 antibodies against SCLC1. The results showed that the immune sera could be diluted > 30 000 and still react with the SCLC1 protein (as compared to the pre-immune serum, which did not contain any significant level of antibodies against SCLC1).

In a Western blot analysis, the SCLC1 protein was also run on an SDS-PAGE gel using the Phast-System (Amersham Pharmacia) and 8-25% gradient gel under reducing 20 conditions. After the electrophoresis had been completed, the SCLC1 protein was diffusion blotted to a nitrocellulose-membrane. After blocking, the membrane was divided and immune serum (dilution 1:10 000 in PBS-0.05%Tween 20) and pre-immune serum, respectively, were added and incubated for two hours at room temp. After washing in PBS-0.05%Tween 20, secondary antibodies (anti rabbit IgG, horseradish-labelled and developed in goat, Sigma) 25 were used to detect rabbit antibodies directed against SCLC1. The result showed that the immune serum detected the immobilized SCLC1 protein efficiently, while no detection was seen using the pre-immune serum.

**Occurrence of sclC gene in strains of subsp. *equi* and subsp. *zooepidemicus***

Chromosomal DNA from different isolates of both subsp. was used as template for a 30 PCR-analysis to study the occurrence of the sclC gene. The PCR primers used were OSCL2:5 (SEQ. ID. NO: 25) and OSCL3:3 (SEQ. ID. NO: 26). The result showed that a similar gene was present in all studied strains of subsp. *equi* and in a majority of strains in subsp. *zooepidemicus*.

In the following Examples 7-10, immunization and challenge of mice were performed as described below.

**Bacterial strain and growth:** *Streptococcus equi* subsp. *equi* was isolated from a horse with strangles. After one passage in a mouse (inoculated intranasally and reisolated 5 from a submandibular gland), the bacteria were stored at -80°C. To prepare an inoculum, stored bacteria were first grown on a blood agar plate, subcultured overnight at 37°C in Todd Hewitt medium containing 1% (w/v) yeast extract. Six ml of this culture were taken into 50 ml of Todd Hewitt /Yeast extract medium (BBL, Becton Dickinson) containing 10% (v/v) equine, i.e. horse, serum (heat inactivated) and incubated at 37°C for 4 hrs. Bacteria grown 10 this way were used in animal challenge experiments.

**Mice and experimental design:** NMRI outbred female mice (Möllegård, Denmark) having a body weight of about 20 g, were individually marked, and kept in groups of four in each cage. Daily, they were weighed and nose impressions onto a BG plate (sheep blood and gentian violet) were taken to quantify bacterial growth. After incubation at 37°C in 5 % CO<sub>2</sub> 15 for 24 hrs, the amount of haemolytic colonies was determined according to a scale of 0-3. A score of 3 means confluent growth of *Streptococcus* subsp *equi* on the whole plate and a score of 0 indicates 0-4 colonies.

**Bronchoalveolar lavage (BAL) and nasal wash (NW) for determination of mucosal IgA responses:** BAL was performed by infusing and withdrawing a solution 20 consisting of 1 ml ice-cold PBS containing protease inhibitor (Complete, Roche) three times into the trachea. NW was performed by rinsing the nasal cavity with 0.5 ml ice-cold PBS containing protease inhibitor. The BAL and NW samples were kept on ice before storage at -80°C.

**Statistical analysis:** Differences between the groups were evaluated by using the 25 Student *t*-test, where *p* < 0.05 was considered to be statistically significant.

**Example 7: Immune response in mice subjected to infection with *Streptococcus* subsp. *equi***

Mice (n=19) were infected with *Streptococcus* subsp. *equi*. Ten µl of bacteria containing 1x10<sup>6</sup> CFU were given intranasally to anaesthetized mice, viz. 5 µl in each nostril. 30 After two weeks, the mice were sacrificed and blood samples were taken. All mice had developed IgG antibodies against FNZN, SFSC1, and EAG4B. The results from eight mice are shown in Figures 1, 2, and 3 for FNZN, SFSC1, and EAG4B, respectively. These ELISA tests were performed according to a standard procedure. Briefly, wells on microtiter plates (Costar) were coated with the respective proteins at a concentration of 10 µg/ml. Serum

dilutions from the mice samples were applied. After washing, rabbit antibodies against mice IgG conjugated with horse radish peroxidase (Dako) were added. To develop the color reaction, OPD tablets (Dako) were added according to instructions from the manufacturer.

**Example 8: Immune response in mice immunized with FNZN, SFSC1, and EAG4B**

Mice (n=24) were immunized subcutaneously on days 0, 7, 14, and 21. Twelve mice were given antigens (FNZN, SFSC1, and EAG4B) and adjuvant and twelve mice were given only adjuvant. The adjuvant used was EtxB provided by Dr Tim R. Hirst, University of Bristol. Each animal was given 12 µg of each antigen and 36 µg of EtxB at each occasion.

10 On day 28, the mice were sacrificed and blood samples were taken. ELISA tests were performed as described above. All mice developed IgG antibodies against FNZN, SFSC1, and EAG4B. Samples from seven mice and a negative control are shown in Figures 4, 5, and 6 for FNZN, SFSC1, and EAG4B, respectively.

**Example 9: Immune response in mice vaccinated with SEC2.16**

15 Mice (n=5) were subcutaneously immunized on days 0, 7, 14, and 21. Each animal was given 12 µg of SEC2.16 in Freund's adjuvant. Serum samples were taken on day 28 and ELISA tests were performed as described above. In Figure 7, the production of IgG antibodies against SEC2.16 is shown.

**Example 10: Subcutaneous immunization with FNZN, SFSC1, and EAG4B followed by challenge with *Streptococcus equi* subsp *equi***

Mice (n=24) were subcutaneously immunized on days 0, 7, 14, and 21. Twelve mice were given antigens (FNZN, SFSC1, and EAG4B) and the adjuvant EtxB and twelve mice were given only adjuvant. Each animal was given 12 µg of each antigen and 36 µg of EtxB at each occasion. On day 28, the mice were infected with *Streptococcus equi* subsp. *equi*. Ten µl of bacteria containing  $1 \times 10^6$  CFU were given intranasally to anaesthetized mice. Animals that lost more than 15% of weight were killed. Mice that had survived or had lost less than 15% of weight are shown in Figure 8. It is evident from Figure 8 that non-vaccinated animals had a lower survival rate than vaccinated animals. On day 3, the p-value was < 0.05. From Figure 9, it is evident that in comparison to non-vaccinated animals, fewer bacteria grew in the noses of vaccinated animals; p-values obtained on days 2, 3, and 7 are 0.03, 0.02, and 0.01, respectively.

**Example 11. Intranasal immunization with FNZN, SFSC1, EAG4B and SEC2.16 followed by challenge with *Streptococcus equi* subsp. *equi***

One group of mice (n=24) was intranasally immunized on days 0, 7, 14, and 21 by administration of the antigens in 40 µl as several small doses in both nostrils. Twelve mice 5 received the antigens FNZN, SFSC1, and EAG4B together with the adjuvant EtxB and twelve mice received only the adjuvant. Each animal was given 12 µg of each antigen together with 36 µg of EtxB at each occasion. On day 28, the mice were anaesthetized and infected with *Streptococcus equi* subsp. *equi* by intranasal administration of 10 µl of bacteria containing  $1 \times 10^6$  CFU (colony forming units).

10 Weight loss versus time is shown in Figure 10; on days 2, 3, 4, 5, and 8, the p-values were 0.04, 0.017, 0.005, 0.003, and 0.002, respectively. Nasal growth versus time is shown in Figure 11; on days 4 to 8, the p-value was <0.001. IgA antibody titers in BAL/NW and sera were determined by ELISA. Briefly, microtiter plates (Costar) were coated with 10 µg/ml of FNZN, SFSC1, and EAG4B. Serial dilutions of BAL/NWs were added to the wells. After 15 washing, goat anti mouse IgA conjugated with horse radish peroxidase (Dako) was added. To develop the color reaction, OPD tablets (Dako) were added. From Figure 12 it is evident that BAL/NW samples from the animals contained IgA. Moreover, IgA against SFSC1 and EAG4B was also found in serum samples from some of the animals (data not shown).

Another group of mice (n=15) was immunized intranasally with SEC2.16 using 20 Matrix (Ref. 20) as adjuvant and a control group was given the Matrix adjuvant alone (n=15). The animals were subjected to challenge infection with subsp. *equi*. The number of animals that died or lost more than 15% weight were different in the two groups; five vaccinated mice and 8 non-vaccinated died (Figure 18). This is not a statistically significant difference but is a 25 tendency implying an effect of SEC2.16. It is likely that it would give a synergistic effect together with any or all of the other antigens mentioned above. Growth of subsp. *equi* in the noses was also studied. More bacteria were isolated from non-vaccinated mice than from vaccinated mice. On day 10, the score was 2.1 and 1.8 (on a scale from 0-3) in these groups, again implying an effect by vaccination with SEC2.16.

**Example 12. Immunization of horses**

30 Horses with no previous history of strangles and with low antibody titers against FNZ, SFS and EAG were selected. Four groups (A-D) of horses with three horses in each group were immunized as follows: A) EAG4B, FNZN and SFSC1 with EtxB as adjuvant, and given both intranasally and subcutaneously; B) As in group A but given only intranasally; C) as in A but without EtxB as adjuvant; and D) Only EtxB, and no antigens, given both

intranasally and subcutaneously. Each dose consisted of 35 µg of each antigen and 100 µg of EtxB in a volume of 2 ml for intranasal and/or subcutaneous immunization. Immunizations were given on days 0, 14, 28 and 42.

Serum samples were taken and analysed as follows: Microtiter wells (Costar) were 5 coated overnight with 100 µl FNZN, SFSC1, or EAG4B, at 10 µg per ml overnight in phosphate buffered saline (PBS). The plates were then blocked with 2% (w/v) bovine serum albumin (BSA) for 1 hour at 37°C. After washing, horse sera were added to wells at a 20-fold dilution, followed by 2-fold serial dilutions. The plates were washed after 1 hour of incubation at 37°C. After washing, detection of antibody binding was performed with 10 antibodies, diluted 1000x, against horse IgG, raised in rabbit and conjugated with HRP (Sigma Chemical Co). Development of a colorimetric reaction was achieved with OPD tablets (Dako, Denmark). Absorbance was determined spectrophotometrically at 492 nm.

In this way, IgG antibodies in sera of the immunized horses were determined. The results are shown in Figure 14. The log dilution of sera required to give an absorbance at a cut 15 off value of 1.0 was calculated for each individual serum sample. Mean values (n=3) with standard errors are shown. Samples taken before (day 0) and after (day 56) immunizations are shown. The horses were divided into four groups A) (white bars): antigens + EtxB given both s.c. and i.n.; B) (striped bars): as group A but immunization only i.n.; C) (gray bars): As in group A but omitting EtxB; D) (black bars): control group given only EtxB both i.n. and s.c.

20

#### **Example 13. Determination in ELISA of antibody titers from horses with or without a history of previous strangles**

Serum samples were taken from horses (n=16) without any previous or present signs of strangles, as well as from horses (n=10) with clinical signs of strangles and positive results 25 from cultivation of *S. equi*. Microtiter wells (Costar) were coated overnight with 100µl FNZN, SFSC1, EAG4B, SEC2.16 or SCLC1 at 10 µg per ml overnight in phosphate buffered saline (PBS). The plates were then blocked with 2% (w/v) BSA for 1 hour at 37°C. After washing, horse sera were added to wells at a 20-fold dilution, followed by 2-fold serial dilutions. The plates were washed after 1 hour incubation at 37°C. Detection of antibody 30 binding was performed after washing, with antibodies, diluted 1000x, against horse IgG, raised in rabbits and conjugated with HRP (Sigma Chemical Co). Development of a colorimetric reaction was achieved with OPD tablets (Dako, Denmark). Absorbance was determined spectrophotometrically at 492 nm. The log<sub>10</sub> dilution of sera, which was required to give an absorbance at a cut-off value of 1.0, was calculated for each individual serum

sample. Mean values of the log10 dilutions, with standard errors, are shown. For comparing normal vs. strangles sera against FNZN, SFSC1, EAG4B, SEC2.16 and SCLC1, the p-values are <0.001, 0.02, <0.0001, 0.02, and <0.0001, respectively.

In Figure 15, the results from a comparison of serum antibody titers from horses with or without strangles are shown. The designation n denotes normal horses without and the designation s denotes horses with strangles. The log dilution of sera required to give an absorbance at a cut off value of 1.0 was calculated for each individual serum sample. Mean values with standard errors are shown. For comparing normal vs. strangles sera against FNZN, SFSC1, EAG4B, SEC2.16, and SCLC1, the p-values are <0.001, 0.02, <0.0001, 0.02 and < 0.0001, respectively.

#### **Example 14. Determination of IgA antibody titers from, immunized horses**

Horses were divided into four groups (A-D) and were immunized as described in Example 12. Antigen-specific IgA in nasal wash samples from immunized horses were determined by an indirect ELISA. Monoclonal antibody K129-3E7 (2 µg/ml) against equine IgA was used to detect bound IgA in the samples followed by use of a goat anti-mouse immunoglobulin HRP conjugate (2 µg/ml). Coating concentrations of antigens were 4 µg/ml. Background values (from no-antigen plates) were subtracted. All samples were analyzed in triplicate. Antigen-specific IgA ELISA data is expressed relative to the total IgA ELISA data in order to correct for nasal secretion dilution during the nasal wash procedure. Total IgA in nasal wash samples was determined by a capture ELISA using two different monoclonal antibodies specific to equine IgA. One was used to coat wells and the other was biotinylated and was detected with the use of a streptavidin-HRP conjugate. The concentrations of monoclonal antibody and streptavidin-HRP were optimized so that none of these was limiting in the assay. Samples were analysed in triplicate. In Figure 16, the Absorbance values obtained for IgA against FNZN and EAG4B are shown.

In Figure 16, IgA antibodies in nasal washings of immunized horses are shown. Mean absorbance values (n=3) in ELISA from groups A-D are shown. The horses were divided into four groups; A) antigens + EtxB given both s.c. and i.n.; B) as group A but immunization only i.n.; C) as in group A but omitting EtxB; D) control group given only EtxB both i.n. and s.c. Light gray bars: samples from day 0 and dark gray bars from day 56.

**Example 15. Determination of kinetics for development of IgG in immunized horses.**

Horses were immunized as described in Example 12. On days 0, 14, 28 and 42, samples were taken on these days and on day 56 as well and analyzed as described elsewhere.

5 The horses were immunized with EAG4B, FNZN and SFSC1 with EtxB, by both intranasal and subcutaneous route.

The antibody titers against EAG4B found in these (three) horses are shown in Figure 17 as a function of time.

**Example 16. Immunogenicity of protein SEC**

10 Mice were immunized intranasally with SEC2.16, with (n=5) or without Matrix (n=5) using 20 µg per mouse of both SEC2.16 and Matrix. Serum samples were taken before and after 4 immunizations, with two-week intervals between. Pre-immune sera did not contain any detectable antibodies against SEC2.16. SEC2.16 without any adjuvant gave significant antibody responses in absence of Matrix. In presence of Matrix even better antibody titers  
15 were obtained as shown in Figure 19.

**Example 17. Comparison of immunization with different antigens followed by challenge infection**

Mice were immunized with proteins FNZN+SFSC1+EAG4B with EtxB (n=8) or with EAG4B with EtxB (n=10) as described in Example 11. Another group was given EtxB  
20 alone as control (n=9). The mice were subjected to challenge with subsp. *equi* and their weight change was monitored. As expected, the control group lost more weight than the groups that were given antigen, which confirms the protective effect of vaccination. The infected mice were left for 14 days. Animals in the group, which was given FNZN+SFSC1+EAG4B with EtxB as adjuvant regained weight from day 10, when average  
25 weight loss was approx. 10%, whereas on day 14, average loss was 7%. On the other hand, mice vaccinated with EAG4B with EtxB as adjuvant, kept loosing weight and on day 14 weight loss in this group was 12.5%. The difference in weight loss on day 14 implies that supplying EAG4B together with other antigens, such as SFSC1 and/or FNZN improves protection. However, the difference is not significant due to small sample size. (No data  
30 shown)

**REFERENCES**

1. Courtney, H. S., Y. Li, J. B. Dale, and D. L. Hasty. 1994. Cloning, sequencing and expression of a fibronectin/fibrinogen-binding protein from group A streptococci. *Infect. Immun.* **62**:3937-3946.
- 5 2. Cue, D., P. E. Dombek, H. Lam, and P.P. Cleary. 1998. *Streptococcus pyogenes* serotype M1 encodes multiple pathways for entry into human epithelial cells. *Infect. Immun.* **66**:4593-4601.
- 10 3. Barnham, M., A. Ljunggren, and M. McIntyre. 1987. Human infection with *Streptococcus zooepidemicus* (Lancefield group C): three case reports. *Epidem. Inf.* **98**:183.-190
- 15 4. Galán, J.E., and J. F. Timoney. 1988. Immunologic and genetic comparison of *Streptococcus equi* isolates from the United States and Europe. *J. Clin. Microbiol.* **26**:1142-1146.
5. Hynes, T. O. 1990. Fibronectins. Springer Verlag, New York.
- 20 6. Engvall, E., E. Ruoslahti, and J. M. Miller. 1978. Affinity of fibronectin to collagen of different genetic types and to fibronogen. *J. Exp. Med.* **147**:1584-1595.
- 25 7. Patti, J. M., Jonsson, H., Guss, B., Switalski, L.M., Wiberg, K., Lindberg, M., and Höök, M. (1992) Molecular characterization and expression of a gene encoding a *Staphylococcus aureus* collagen adhesin. *J. Biol. Chem.* **267**:4766-4772.
8. Jonsson, H., Lindmark, H., and Guss, B. (1995) A protein G related cell surface protein in *Streptococcus zooepidemicus*. *Infect Immun* **63**:2968-2975.
- 30 9. Lindmark, H., Jacobsson, K., Frykberg, L., and Guss, B. (1996) Fibronectin-binding protein of *Streptococcus equi* subspecies *zooepidemicus*. *Infect Immun* **64**:3993-3999.
10. Jacobsson, K., Jonsson, H., Lindmark, H., Guss, B., Lindberg, M., and Frykberg, L. (1997) Shot-gun phage display mapping of two streptococcal cell-surface proteins. *Microbiol Res.* **152**:1-8.

11. Lindmark, H., Jonsson, P., Olsson-Engvall, E., and Guss, B. (1999) Pulsed-field gel electrophoresis and distribution of the genes *zag* and *fnz* in isolates of *Streptococcus equi*. *Res Vet Sci.* 66:93-99.

5

12. Lindmark, H., and Guss, B. (1999) SFS, a novel fibronectin-binding protein from *Streptococcus equi*, inhibits the binding between fibronectin and collagen. *Infect. Immun.* 67: 2383-2388.

10 13. Lindmark, H. (1999) Characterization of adhesive extracellular proteins from *Streptococcus equi*. (Doctoral thesis) *Acta Universitatis Agriculturae Sueciae, Agraria* 139. ISBN 91-576-5488-3

14. Lindmark, H., Nilsson, M., and Guss, B. (2001) Comparison of the fibronectin-15 binding protein FNE from *Streptococcus equi* subspecies *equi* with FNZ from *S. equi* subspecies *zooepidemicus* reveals a major and conserved difference. 69: 3159-3163.

15. Schneewind, O., Fowler, A. and Faull, K.F. (1995) Structure of the cell wall anchor of surface proteins in *Staphylococcus aureus*. *Science* 268:103-106.

20

16. Anton Mayr et al. *Handbuch der Schutzimpfungen in der Tiermedizin*.3.3.4. p. 196-200. Verlag Paul Parey. Berlin and Hamburg. 11984

17. Elson, C.O., and M.T. Dertzbaugh. 1999. p. 817-838, *Nucosal immunology*, 2<sup>nd</sup> ed.

25 Of Academic Press, New York, N.Y.

18. Winner et al. 1991. *Infect. Immun.* 59:997-982

19. Ogra et al. *Clinical Microbiology Reviews*, Apr. 2002, p. 430-445.

30

20. B. Morein and Karin Lövgren Bengtsson 1998, 76:295-299. *Immunology and Cellbiology*

21. **Hanski, E., and M.G. Caparon.** 1992. Protein F, a fibronectin-binding protein, is an adhesin of the group A streptococcus *Streptococcus pyogenes*. Proc. Natl. Acad. Sci. USA 89:6172-6176.

5 22. **Hanski, E., P.A. Horwitz, and M.G. Caparon.** 1992. Expression of protein F, the Fibronectin-binding protein of *Streptococcus pyogenes* JRS4, in heterologous streptococcal and enterococcal strains promotes their adherence to respiratory epithelial cells. Infect. Immun. 60:5119-5125.

10 23. **Jadoun, J., V. Ozeri, E. Burstein, E. Skutelsky, E. Hanski, and S. Sela.** 1998. Protein F1 required for efficient entry of *Streptococcus pyogenes* into epithelial cells. J. Infect. Dis. 178:147-158.

24. **Molinri, G., S. R. Talay, P. Valentin-Weigand, M. Rohde, and G.S. Chhatwal.** 1997.

15 The fibronectin-binding protein of *Streptococcus pyogenes* SfbI, is involved in the internalization of group A streptococci by epithelial cells. Infect. Immun. 65:1357-1363.

25. **Lannergård, J., Frykberg, L. and Guss B.** (2003) CNE, a collagen-binding protein of *Streptococcus equi*. FEMS Microbiol. Lett. 222:69-74

20

26. WO 92/07002

27. WO 00/37496

**CLAIMS**

- 5 1. An antigenic composition comprising at least one antigen, wherein said at least one antigen comprises at least part of a protein of *Streptococcus equi* subsp *equi*, and said at least part of said protein comprises at least one antigenic epitope or antigenic determinant of *Streptococcus equi*, and wherein said protein is comprised of EAG comprising an amino acid sequence according to SEQ ID NO: 1, or an analog thereof.
- 10
2. The antigenic composition of claim 1, which comprises at least one further antigen that comprises at least part of a fibronectin-binding protein of *Streptococcus equi*, said at least part of said protein comprising at least one antigenic epitope or antigenic determinant of *Streptococcus equi*, and wherein said protein is selected from the group consisting of FNZ comprising an amino acid sequence according to SEQ ID NO: 2 and SFS comprising an amino acid sequence according to SEQ ID NO: 3, or an analog thereof.
- 15
- 20 3. The antigenic composition of claim 1 or 2, which comprises at least one further antigen that comprises at least part of a protein of *Streptococcus equi* and said at least part of said protein comprises an antigenic epitope or an antigenic determinant of *Streptococcus equi*, and wherein said protein is comprised of SEC comprising an amino acid sequence according to SEQ ID NO: 4, or an analog thereof.
- 25
4. The antigenic composition of claim 1, 2, or 3 which comprises an N-terminal fragment of a protein selected from the group consisting of EAG and FNZ.
5. The antigenic composition of claim 3, wherein said antigens are comprised of at least part of EAG, FNZ, SFS, and SEC, optionally, said at least part of EAG and FNZ being an N-terminal part thereof, said at least part of SFS being a C-terminal part of SFS, and said at least part of SEC being a collagen-binding part of SEC.
- 30

6. A vaccine composition for protecting non-human mammals against infection of *Streptococcus equi*, which comprises the antigenic composition of claim 1 as immunizing component and a pharmaceutically acceptable carrier.
- 5 7. The vaccine composition of claim 6, which comprises the antigenic composition of any one of claims 2-5 as immunizing component.
8. The vaccine composition of claim 6 or 7, which further comprises an adjuvant.
- 10 9. The vaccine composition of any one of claims 6-8, which is a vaccine that protects susceptible mammals, suitably horses, against strangles caused by *Streptococcus equi* subsp. *equi*.
- 15 10. The vaccine composition of claim 7, which is provided in a physiologically administrable form and suitably is administrable by subcutaneous or intranasal inoculation.
- 20 11. The vaccine composition of claim 9 or 10, which stimulates serum, mucosal and/or bronchial lavage antibody responses directed to *Streptococcus equi* antigens in mammals susceptible to *Streptococcus equi*, suitably horses.
12. A method for producing an antigen of an antigenic composition of any one of claims 1-5, which method comprises
  - (a) providing a DNA fragment encoding said antigen and introducing said fragment into an expression vector;
  - (b) introducing said vector, which contains said DNA fragment, into a compatible host cell;
  - (c) culturing said host cell provided in step (b) under conditions required for expression of the product encoded by said DNA fragment; and
  - (d) isolating the expressed product from the cultured host cell, and, optionally,
  - (e) purifying the isolated product from step (d) by affinity chromatography or other chromatographic methods known in the art.
- 25
- 30

13. A method for preparation of a vaccine according to any one of claims 6-11, which vaccine contains as immunizing component, an antigenic composition of any one of claims 1-5, said method comprising mixing said antigenic composition and a pharmaceutically acceptable carrier.

5

14. Use of an antigenic composition of any one of claims 1-5 in the preparation of a vaccine protecting against *S. equi* infection inclusive of strangles caused by subsp. *equi* infection in horses.

10

15. A method for the production of an antiserum, said method comprising administering an antigenic preparation of any one of claims 1-5 to an animal host to produce antibodies in said animal host and recovering antiserum containing said antibodies produced in said animal host.

15

16. A method of prophylactic or therapeutic treatment of *S. equi* infection in non-human mammals, suitably horses, comprising administering to said mammal an immunologically effective amount of a vaccine of any one of claim 6-11 or an antiserum produced according to claim 15.

20

17. A method for protecting horses against *Streptococcus equi* infection, which comprises inoculating a horse subcutaneously or intranasally with a vaccine of any one of claims 6-11 to induce an immune response against *Streptococcus equi* in said horse.

25

18. The method of claim 17, wherein an immune response in the form of IgG and/or IgA and/or IgM antibodies in the nasopharyngeal mucus is induced in said horse.

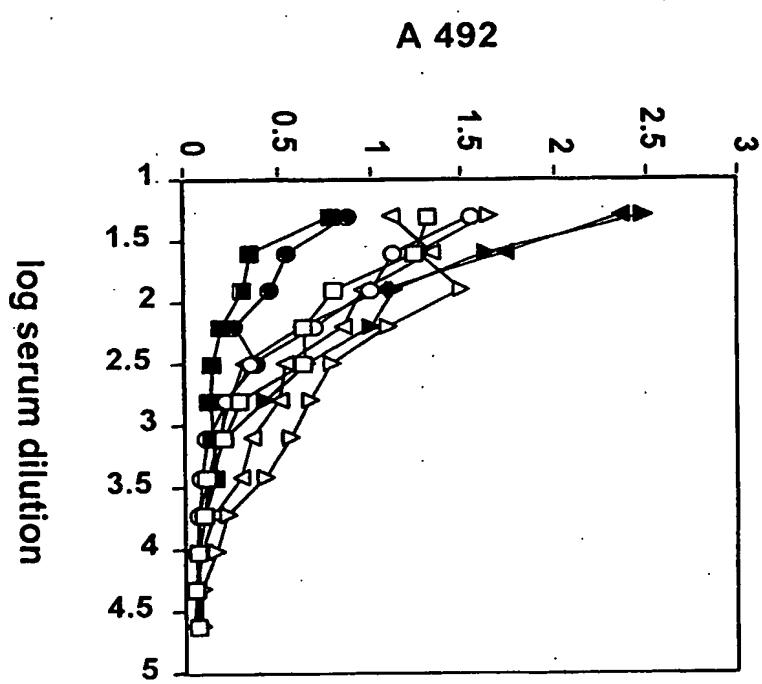
19. Monoclonal antibodies against antigen(s) of the composition of any of claims 1-4 to be used prophylactically or therapeutically against strangles.

30

20. The antigenic composition of any one of claims 1-5, which comprises at least part of a protein designated Scl and comprising an amino acid sequence according to SEQ ID. NO:23 or a fragment thereof, suitably a fragment designated SCL C1 and

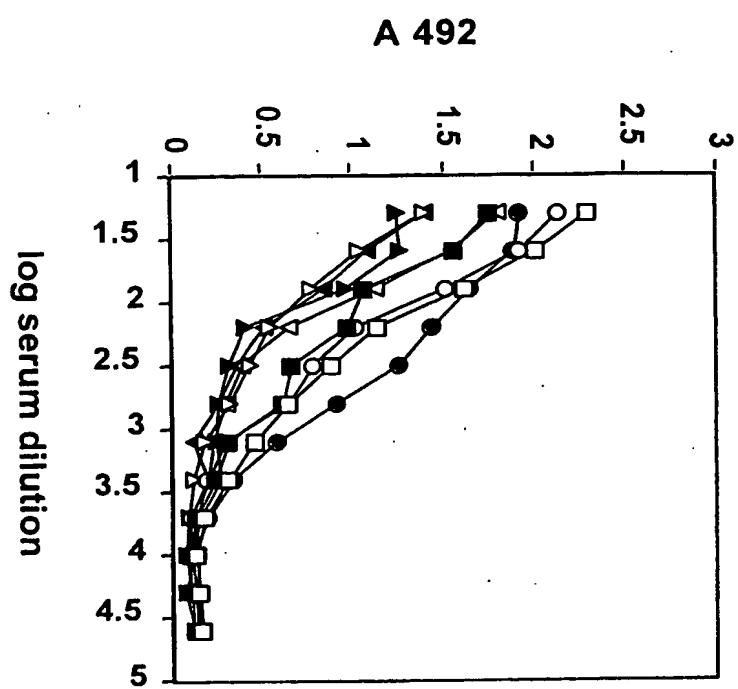
comprising an amino acid sequence according to SEQ. ID. NO:27, or an analog thereof.

Fig 1



IgG antibodies against FNZN, 2 weeks  
after intranasal inoculation with  
**Streptococcus equi subsp. equi**

Fig 2



IgG antibodies against SFSC1, 2 weeks  
after intranasal inoculation with  
*Streptococcus equi* subsp. *equi*

Fig 3

IgG antibodies against EAG4B, 2 weeks  
after intranasal inoculation with  
*Streptococcus equi* subsp. *equi*

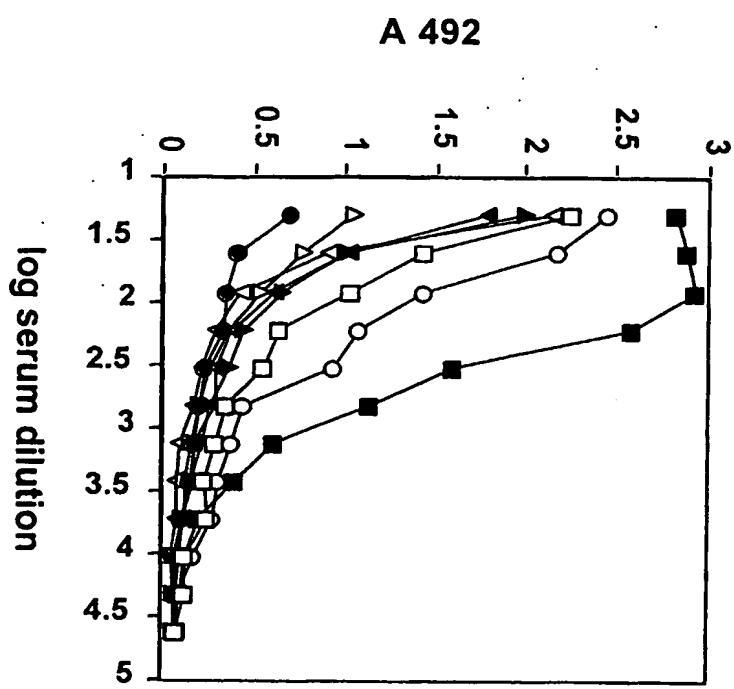
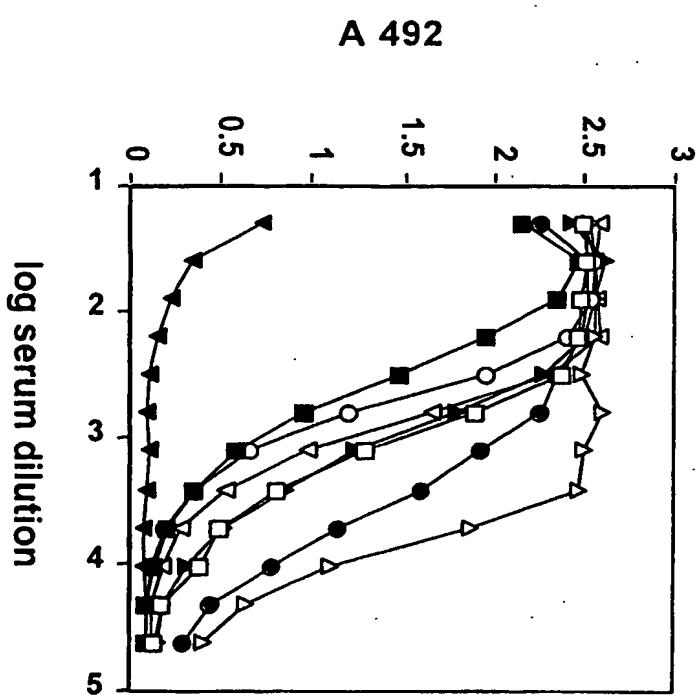
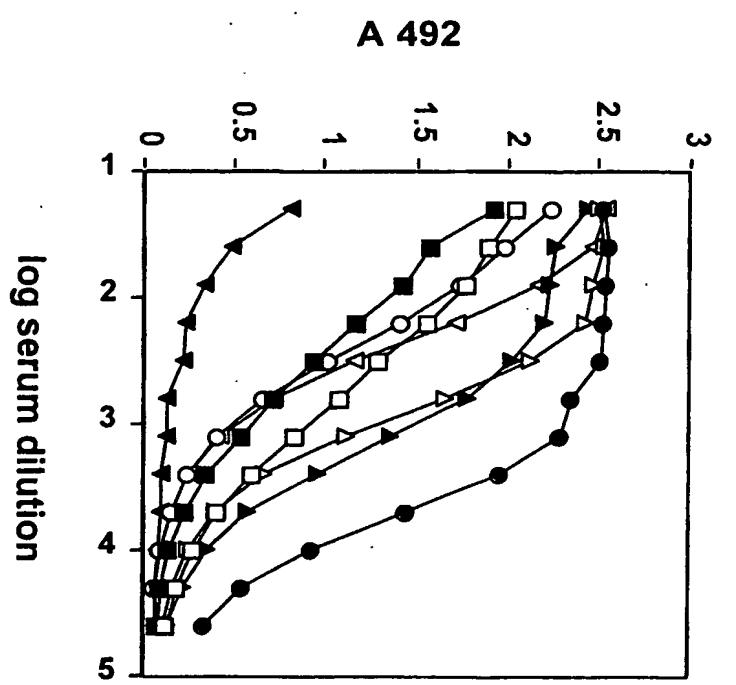


Fig 4



IgG antibodies against FNZN  
after subcutaneous immunization  
with FNZN, SFSC1 or EAG4B

Fig 5



IgG antibodies against SFSC1  
after subcutaneous immunization  
with FNZN, SFSC1  $\circ$  EAG4B

Fig 6

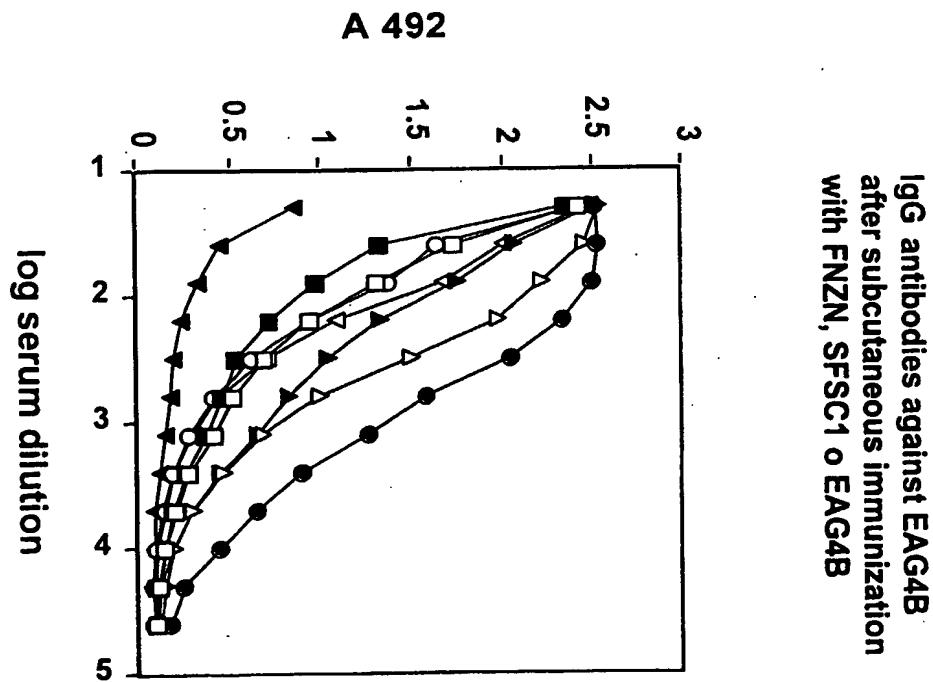
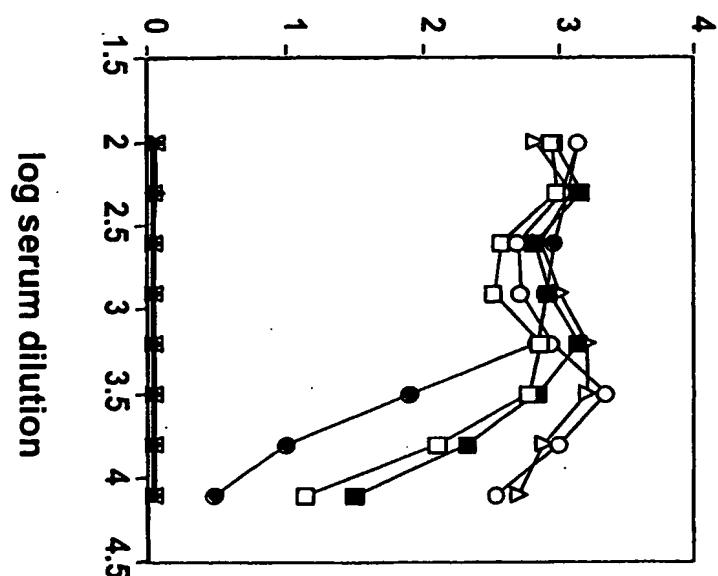


Fig 7

A 492



IgG antibodies against SEC 2.16 after  
subcutaneus immunization with SEC 2.16

Fig 8

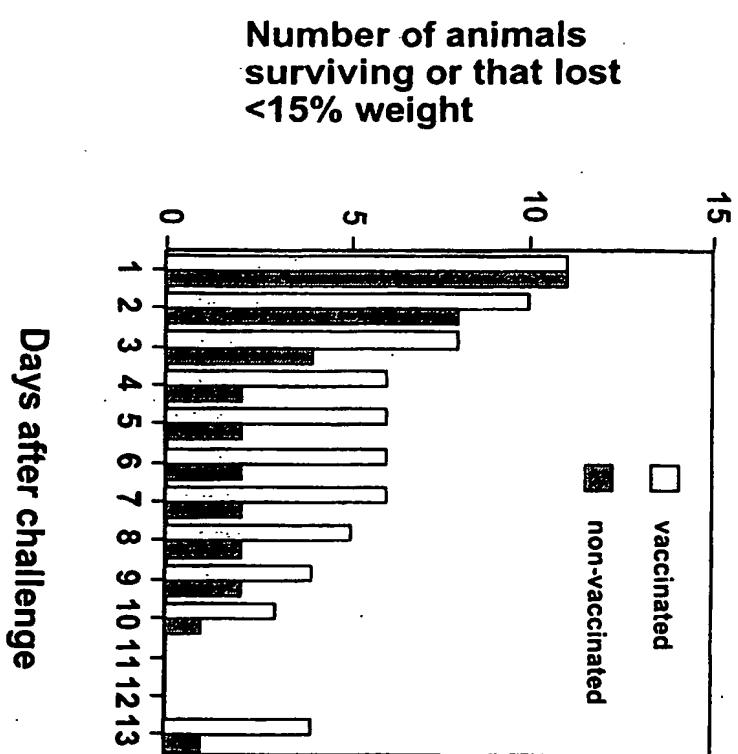


Fig 9

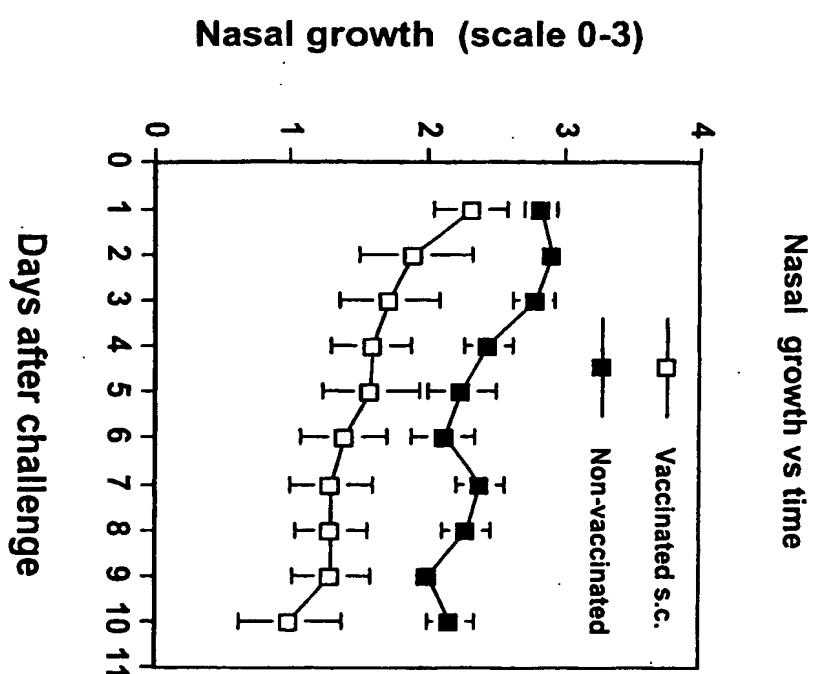


Fig 10

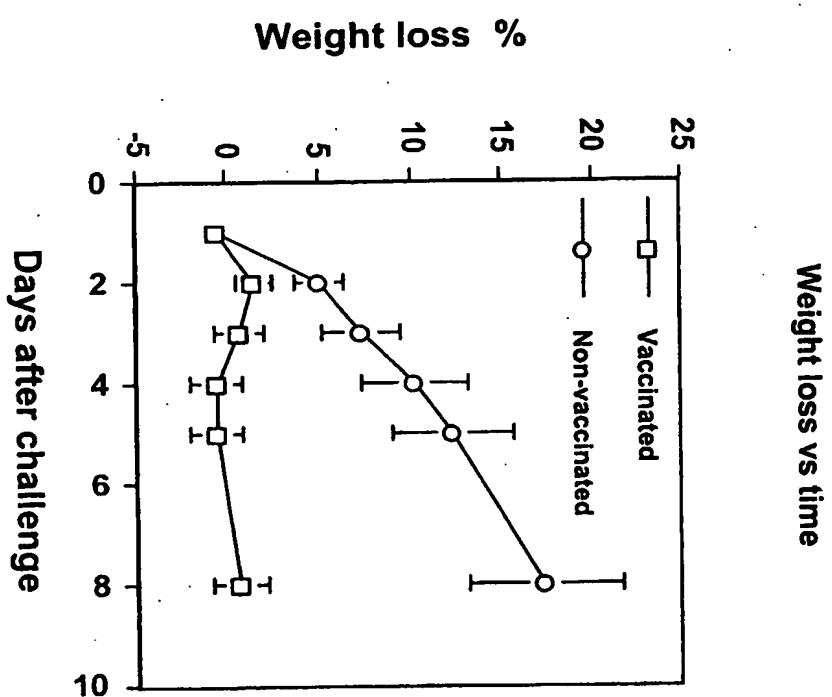


Fig 11

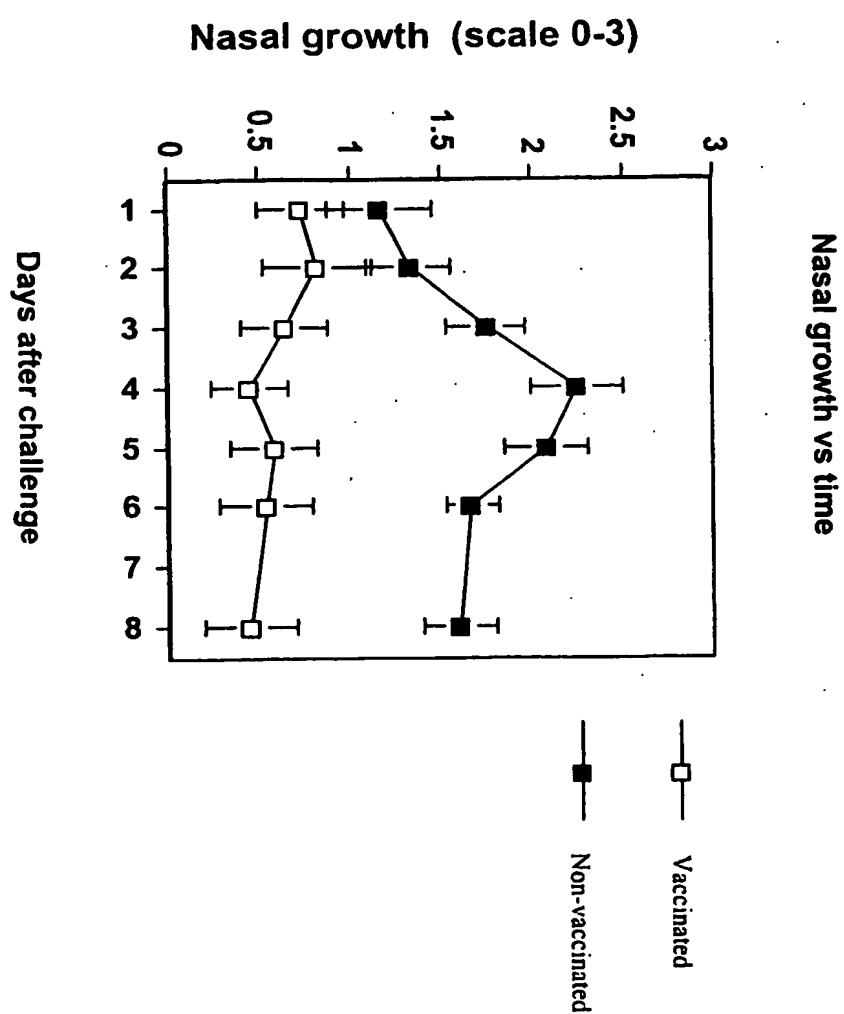


Fig 12

BAL/NW IgA antibodies against  
SFSC1 after intranasal immunization  
with FNZN, SFSC1 and EAG4B

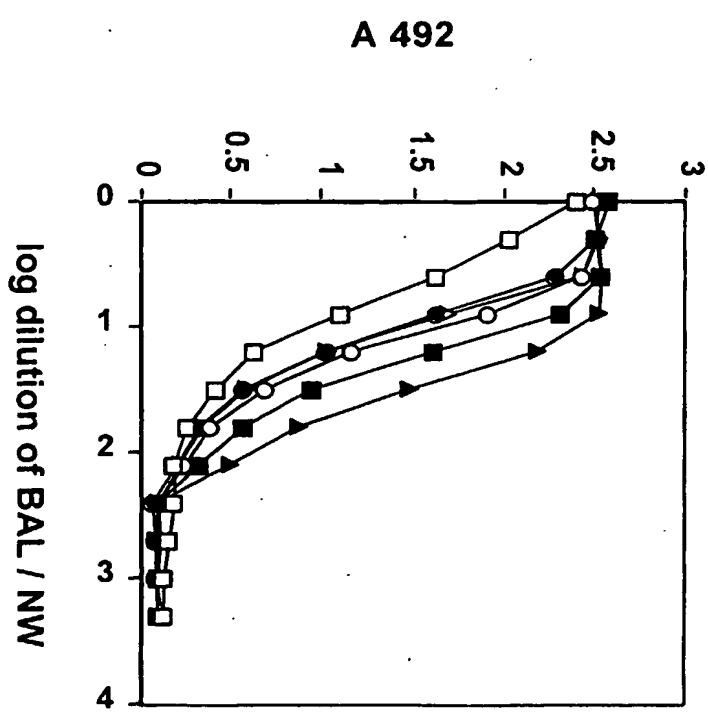


Fig 13

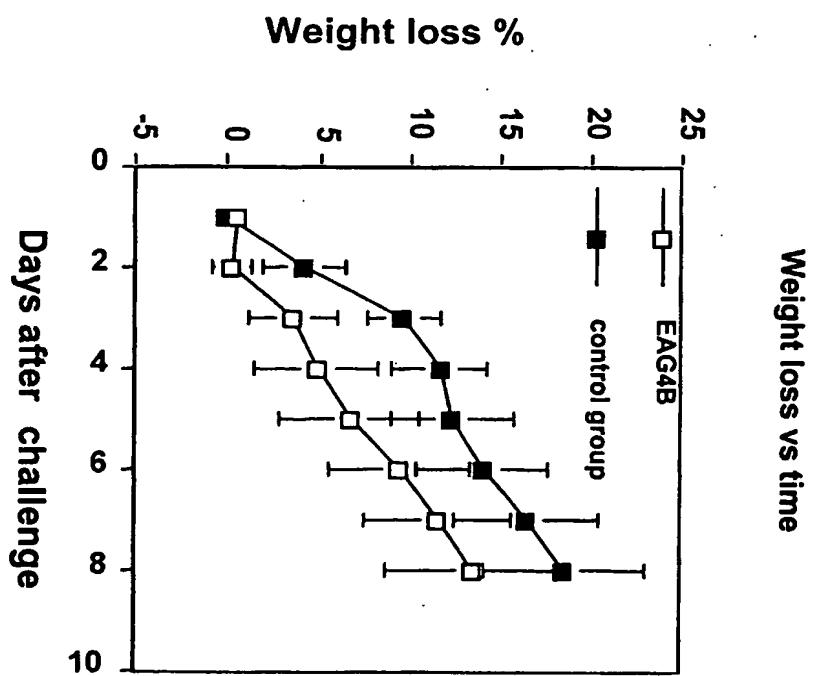
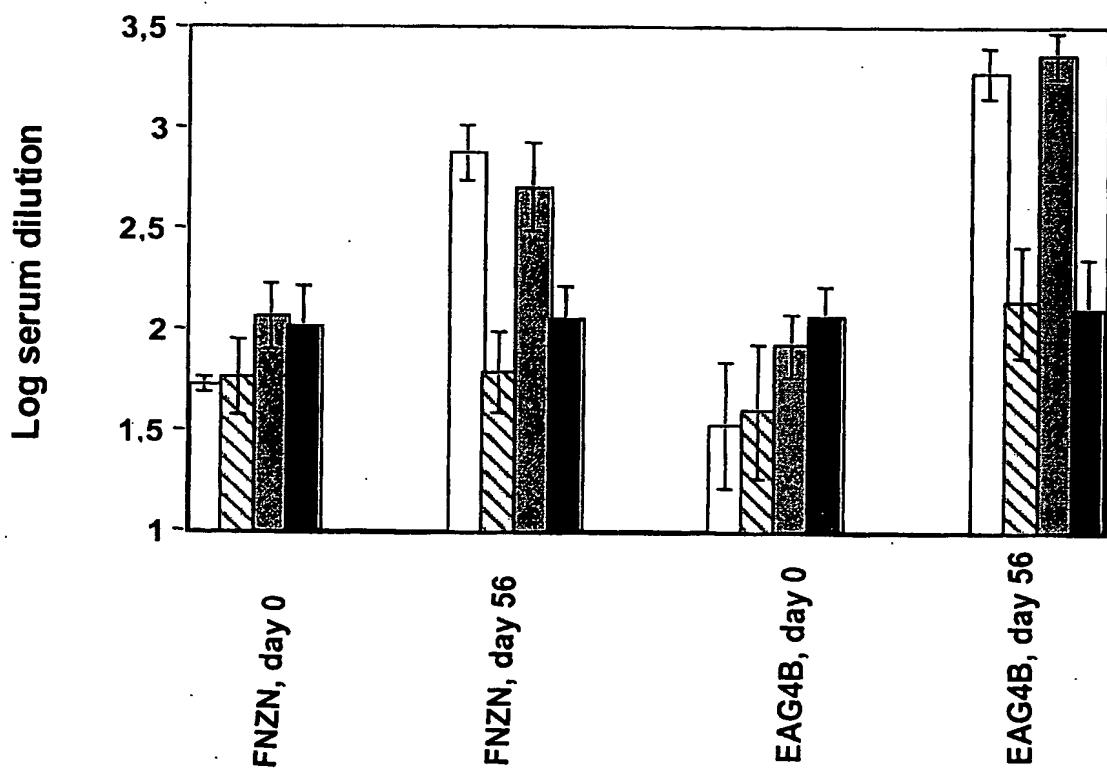
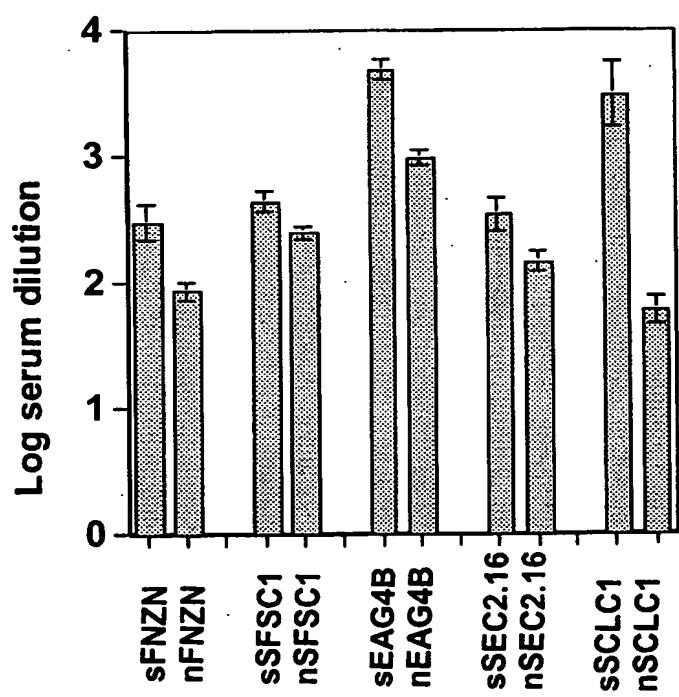
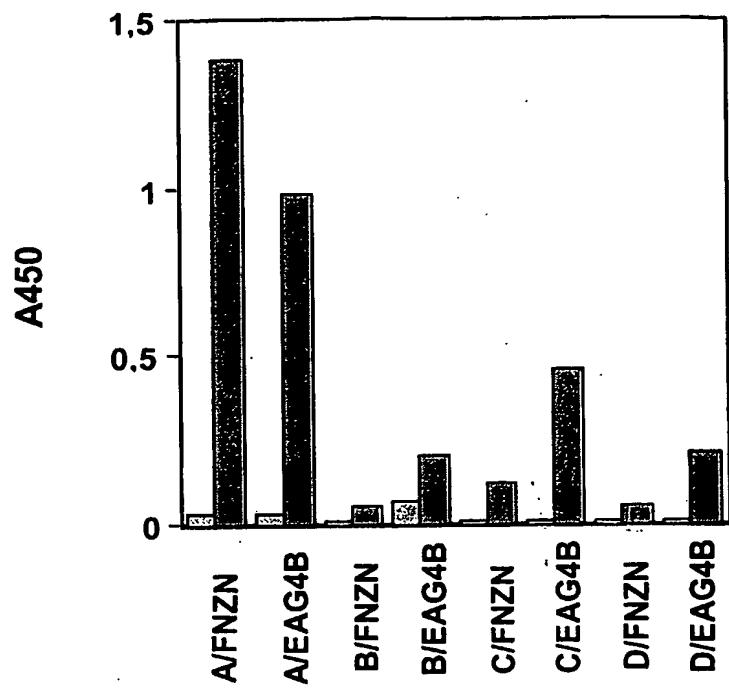
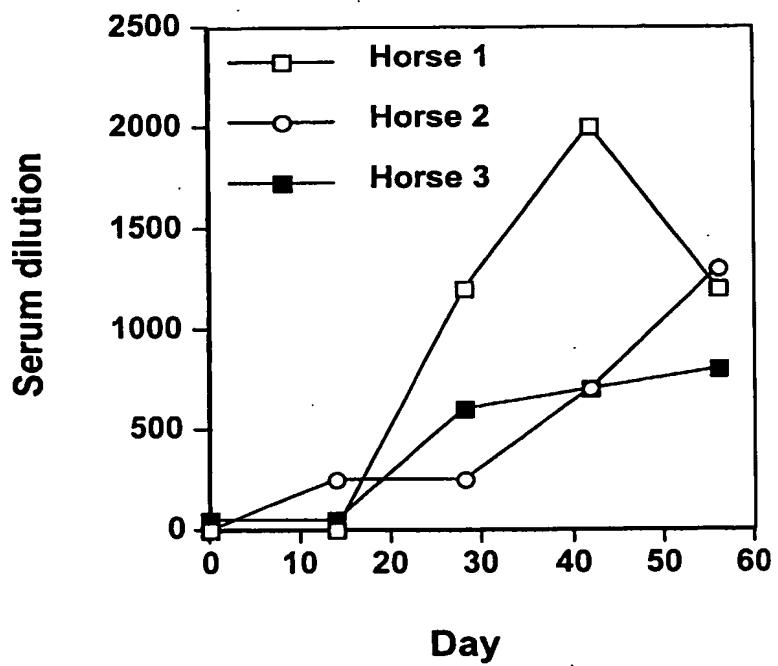


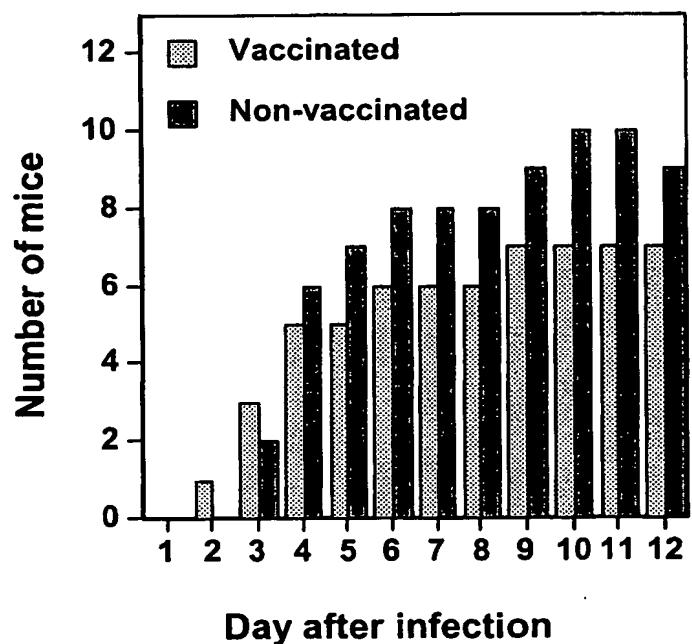
Figure 14



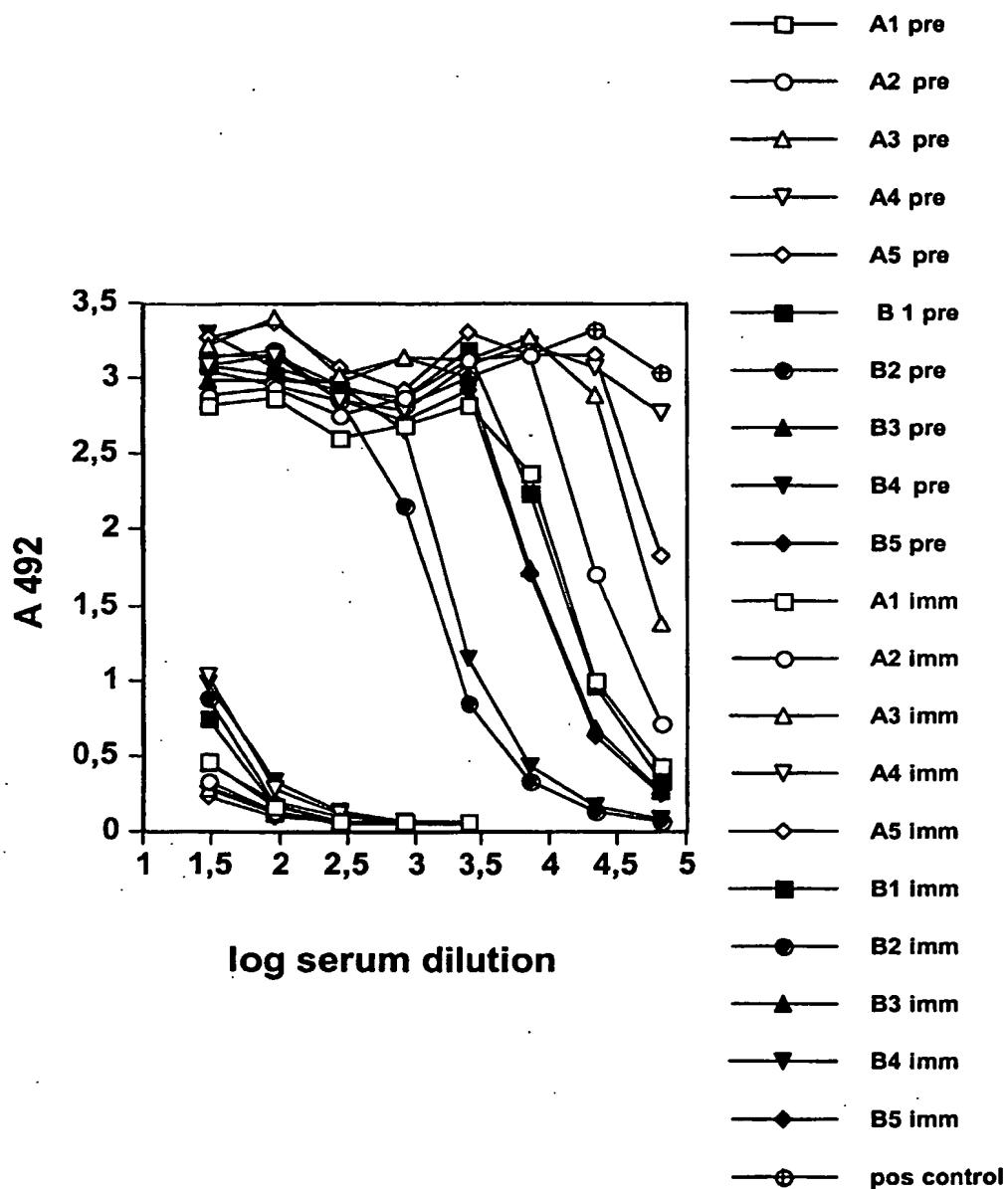
**Figure 15**

**Figure 16**

**Figure 17**

**Figure 18**

**Figure 19**



110087400US.ST25  
SEQUENCE LISTING

<110> Guss et al., Bengt

<120> Immunization of non-human mammals against *Streptococcus equi*

<130> 110087401 PC

<150> US 60/417,660

<151> 2002-10-11

<160> 27

<170> PatentIn version 3.1

<210> 1

<211> 180

<212> PRT

<213> *Streptococcus equi*

<400> 1

Met Ala Leu Asp Ala Thr Thr Val Leu Glu Pro Thr Thr Ala Phe Ile  
1 5 10 15

Arg Glu Ala Val Arg Glu Ile Asn Gln Leu Ser Asp Asp Tyr Ala Asp  
20 25 30

Asn Gln Glu Leu Gln Ala Val Leu Ala Asn Ala Gly Val Glu Ala Leu  
35 40 45

Ala Ala Asp Thr Val Asp Gln Ala Lys Ala Ala Leu Asp Lys Ala Lys  
50 55 60

Ala Ala Val Ala Gly Val Gln Leu Asp Glu Ala Arg Arg Glu Ala Tyr  
65 70 75 80

Arg Thr Ile Asn Ala Leu Ser Asp Gln His Lys Ser Asp Gln Lys Val  
85 90 95

Gln Leu Ala Leu Val Ala Ala Ala Lys Val Ala Asp Ala Ala Ser  
100 105 110

110087400US.ST25

Val Asp Gln Val Asn Ala Ala Ile Asn Asp Ala His Thr Ala Ile Ala  
115 120 125

Asp Ile Thr Gly Ala Ala Leu Leu Glu Ala Lys Glu Ala Ala Ile Asn  
130 135 140

Glu Leu Lys Gln Tyr Gly Ile Ser Asp Tyr Tyr Val Thr Leu Ile Asn  
145 150 155 160

Lys Ala Lys Thr Val Glu Gly Val Asn Ala Leu Lys Ala Lys Ile Leu  
165 170 175

Ser Ala Leu Pro  
180

<210> 2

<211> 597

<212> PRT

<213> *Streptococcus equi*

<400> 2

Met Lys Thr Lys Ser Phe Arg Lys Val Leu Thr Thr Ser Ala Thr Cys  
1 5 10 15

Ile Val Leu Ala Thr Ser Phe Ala Gly Gly Thr Leu Arg Val Trp Ala  
20 25 30

Glu Gln Leu Tyr Tyr Gly Trp Asn Asp Gly Thr Arg Gln Ser Ser Pro  
35 40 45

Tyr Phe Leu Tyr Val Ser Pro Lys Asn Ala Pro Lys Arg Glu Leu Lys  
50 55 60

Asp Glu Tyr Val Val Tyr Cys Phe Asn Lys Lys Leu Tyr Trp Pro Asp  
65 70 75 80

Gln Trp Glu Ser Ile Tyr Ser Asn Phe Asn Asp Ile Arg Ser Pro Tyr  
85 90 95

Asn Asp Leu Pro Val Tyr Glu Lys Lys Leu Gly Tyr Asp Gly Ile Phe  
100 105 110

Lys Gln Tyr Ala Pro Asp Tyr Lys Lys Asp Ile Ser Asp Ile Ala Ser  
115 120 125

Ala Leu Val Ala Val Leu Ser Asn Gly Tyr Pro Thr Asn Lys Ser Gln  
130 135 140

110087400US.ST25

Leu Ser Thr Ser Tyr His Leu Asn Asn Asp Ser Ser Arg Lys Val Thr  
145 150 155 160

Gln Leu Ala Ile Trp Tyr Phe Ser Asp Ser Leu Thr Lys Glu Tyr Leu  
165 170 175

Lys Asp Thr Gly Gly Tyr Asn Leu Asn Asp Met Glu Lys Lys Ala Leu  
180 185 190

Asp Phe Leu Ile Ser Lys Gly Glu Asp Ser Lys Leu Lys Ser Glu Gln  
195 200 205

Ser Asn Tyr Ser Leu Asp Ile Tyr Val Tyr Gln Ser Gly Gly His Asp  
210 215 220

His Met Lys Asp Tyr Gln Asn Leu Leu Gly Ser Thr Leu Ile Pro Lys  
225 230 235 240

Glu Pro Leu Lys Pro Gln Leu Gly Gly Phe Ser Gly His Asn Gly Asn  
245 250 255

Gly Leu Ser Gly Leu Glu Gly Ser Ser Gly Ser Gln Glu Thr Asn  
260 265 270

Glu Asp Gly Lys Lys Gly Leu Ile Gly Phe His Gly Gly Leu Ser Gly  
275 280 285

Ser Glu Gly Lys Arg Asp Pro Leu Pro Gly Leu Lys Gly Glu Ala Gly  
290 295 300

Ala Pro Asp Thr Pro Gln Lys Pro Asn Asp Pro Leu Gln Gly Leu Glu  
305 310 315 320

Gly Gly Asn Ser Pro Ile Val Glu Gln Asn Tyr Gly Ser Thr Glu Gly  
325 330 335

Tyr His Gly Gln Ser Gly Ile Leu Glu Glu Thr Glu Asp Thr Asn Pro  
340 345 350

Pro Gly Ile Ile Leu Gly Gly Ser Gly Asn Val Glu Thr His Glu Asp  
355 360 365

Thr Arg Asn Pro His Leu Met Gly Ile Gly Gly Leu Ala Gly Glu  
370 375 380

Ser Gly Glu Thr Thr Pro Lys Pro Gly Gln Thr Gly Gly Gln Gly Pro  
385 390 395 400

Val Ile Glu Thr Thr Glu Asp Thr Gln Lys Gly Met Ser Gly Gln Ser  
405 410 415

110087400US.ST25

Gly Gly Thr Ile Glu Ser Glu Asn Thr Lys Lys Pro Glu Val Met Ile  
420 425 430

Gly Gly Gln Gly Gln Thr Ile Glu Thr Thr Glu Asp Thr Gln Lys Gly  
435 440 445

Met Ser Gly Gln Ser Gly Gly Thr Ile Glu Ser Glu Asp Thr Lys Lys  
450 455 460

Pro Glu Val Met Ile Gly Gly Gln Gln Ile Ile Asp Phe Ser Glu  
465 470 475 480

Asn Thr Gln Ser Gly Met Ser Gly Gln Ser Gly Asp Thr Thr Val Ile  
485 490 495

Glu Asp Thr Lys Lys Ser Glu Ile Ile Gly Gly Gln Gln Ile  
500 505 510

Ile Asp Phe Ser Glu Asp Thr Gln Pro Gly Met Ser Gly Gln Ser Gly  
515 520 525

Gly Thr Thr Ile Val Glu Asp Thr Lys Lys Pro Thr Pro Lys Pro Lys  
530 535 540

Pro Ala Pro Ala Pro Ile Val Asn Asp Glu Lys Pro Asn Lys Gly Thr  
545 550 555 560

His Leu Pro Gln Thr Ser Asp Met Lys Gln Leu Thr Leu Ser Ile Ile  
565 570 575

Gly Ala Met Ser Met Leu Leu Val Leu Cys Leu Ser Leu Phe Lys Arg  
580 585 590

Pro Ser Lys Lys Asp  
595

<210> 3

<211> 371

<212> PRT

<213> *Streptococcus equi*

<400> 3

Met Arg Lys Thr Glu Gly Arg Phe Arg Thr Trp Lys Ser Lys Lys Gln  
1 5 10 15

Trp Leu Phe Ala Gly Ala Val Val Thr Ser Leu Leu Leu Gly Ala Ala  
20 25 30

110087400US.ST25

Leu Val Phe Gly Gly Leu Leu Gly Ser Leu Gly Gly Ser Ser His Gln  
35 40 45

Ala Arg Pro Lys Glu Gln Pro Val Ser Ser Ile Gly Asp Asp Asp Lys  
50 55 60

Ser His Lys Ser Ser Ser Asp Ser Met Val Ser Arg Pro Pro Lys Lys  
65 70 75 80

Asp Asn Leu Gln Pro Lys Pro Ser Asp Gln Pro Thr Asn His Gln His  
85 90 95

Gln Ala Thr Ser Pro Ser Gln Pro Thr Ala Lys Ser Ser Gly His His  
100 105 110

Gly Asn Gln Pro Gln Ser Leu Ser Val Asn Ser Gln Gly Asn Ser Ser  
115 120 125

Gly Gln Ala Ser Glu Pro Gln Ala Ile Pro Asn Gln Gly Pro Ser Gln  
130 135 140

Pro Leu Gly Leu Arg Gly Gly Asn Ser Ser Gly Ser Gly His His His  
145 150 155 160

Gln Pro Gln Gly Lys Pro Gln His Leu Asp Leu Gly Lys Asp Asn Ser  
165 170 175

Ser Pro Gln Pro Gln Pro Lys Pro Gln Gly Asn Ser Pro Lys Leu Pro  
180 185 190

Glu Lys Gly Leu Asn Gly Glu Asn Gln Lys Glu Pro Gln Gly Glu  
195 200 205

Arg Gly Glu Ala Gly Pro Pro Leu Ser Gly Leu Ser Gly Asn Asn Gln  
210 215 220

Gly Arg Pro Ser Leu Pro Gly Leu Asn Gly Glu Asn Gln Lys Glu Pro  
225 230 235 240

Glu Gln Gly Glu Arg Gly Glu Ala Gly Pro Pro Ser Thr Pro Asn Leu  
245 250 255

Glu Gly Asn Asn Arg Lys Asn Pro Leu Lys Gly Leu Asp Gly Glu Asn  
260 265 270

Lys Pro Lys Glu Asp Leu Asp Gly Lys Gly Leu Ser Gly Glu Asn Asp  
275 280 285

Glu Ser Pro Lys Leu Lys Asp Glu His Pro Tyr Asn His Gly Arg Arg  
290 295 300

110087400US.ST25

Asp Gly Tyr Arg Val Gly Tyr Glu Asp Gly Tyr Gly Gly Lys Lys His  
305 310 315 320

Lys Gly Asp Tyr Pro Lys Arg Phe Asp Glu Ser Ser Pro Lys Glu Tyr  
325 330 335

Asn Asp Tyr Ser Gln Gly Tyr Asn Asp Asn Tyr Gly Asn Gly Tyr Leu  
340 345 350

Asp Gly Leu Ala Asp Arg Gly Gly Lys Arg Gly Tyr Gly Tyr Ser Tyr  
355 360 365

Asn Pro Asp  
370

<210> 4

<211> 657

<212> PRT

<213> *Streptococcus equi*

<400> 4

Leu Lys Gln Leu Thr Lys Ile Val Ser Val Val Leu Leu Leu Val Phe  
1 5 10 15

Thr Leu Ser Ala Ser Leu His Lys Val Arg Ala Thr Asn Leu Ser Asp  
20 25 30

Asn Ile Thr Ser Leu Thr Val Ala Ser Ser Ser Leu Arg Asp Gly Glu  
35 40 45

Arg Thr Thr Val Lys Val Ala Phe Asp Asp Lys Lys Gln Lys Ile Lys  
50 55 60

Ala Gly Asp Thr Ile Glu Val Thr Trp Pro Thr Ser Gly Asn Val Tyr  
65 70 75 80

Ile Gln Gly Phe Asn Lys Thr Ile Pro Leu Asn Ile Arg Gly Val Asp  
85 90 95

Val Gly Thr Leu Glu Val Thr Leu Asp Lys Ala Val Phe Thr Phe Asn  
100 105 110

Gln Asn Ile Glu Thr Met His Asp Val Ser Gly Trp Gly Glu Phe Asp  
115 120 125

Ile Thr Val Arg Asn Val Thr Gln Thr Thr Ala Glu Thr Ser Gly Thr  
130 135 140

110087400US.ST25

Thr Thr Val Lys Val Gly Asn Arg Thr Ala Thr Ile Thr Val Thr Lys  
145 150 155 160

Pro Glu Ala Gly Thr Gly Thr Ser Ser Phe Tyr Tyr Lys Thr Gly Asp  
165 170 175

Met Gln Pro Asn Asp Thr Glu Arg Val Arg Trp Phe Leu Leu Ile Asn  
180 185 190

Asn Asn Lys Glu Trp Val Ala Asn Thr Val Thr Val Glu Asp Asp Ile  
195 200 205

Gln Gly Gln Thr Leu Asp Met Ser Ser Phe Asp Ile Thr Val Ser  
210 215 220

Gly Tyr Arg Asn Glu Arg Phe Val Gly Glu Asn Ala Leu Thr Glu Phe  
225 230 235 240

His Thr Thr Phe Pro Asn Ser Val Ile Thr Ala Thr Asp Asn His Ile  
245 250 255

Ser Val Arg Leu Asp Gln Tyr Asp Ala Ser Gln Asn Thr Val Asn Ile  
260 265 270

Ala Tyr Lys Thr Lys Ile Thr Asp Phe Asp Gln Lys Glu Phe Ala Asn  
275 280 285

Asn Ser Lys Ile Trp Tyr Gln Ile Leu Tyr Lys Asp Gln Val Ser Gly  
290 295 300

Gln Glu Ser Asn His Gln Val Ala Asn Ile Asn Ala Asn Gly Gly Val  
305 310 315 320

Asp Gly Ser Arg Tyr Thr Ser Phe Thr Val Lys Lys Ile Trp Asn Asp  
325 330 335

Lys Glu Asn Gln Asp Gly Lys Arg Pro Lys Thr Ile Thr Val Gln Leu  
340 345 350

Tyr Ala Asn Asp Gln Lys Val Asn Asp Lys Thr Ile Glu Leu Ser Asp  
355 360 365

Thr Asn Ser Trp Gln Ala Ser Phe Gly Lys Leu Asp Lys Tyr Asp Ser  
370 375 380

Gln Asn Gln Lys Ile Thr Tyr Ser Val Lys Glu Val Met Val Pro Val  
385 390 395 400

Gly Tyr Gln Ser Gln Val Glu Gly Asp Ser Gly Val Gly Phe Thr Ile  
405 410 415

110087400US.ST25

Thr Asn Thr Tyr Thr Pro Glu Val Ile Ser Ile Thr Gly Gln Lys Thr  
420 425 430

Trp Asp Asp Arg Glu Asn Gln Asp Gly Lys Arg Pro Lys Glu Ile Thr  
435 440 445

Val Arg Leu Leu Ala Asn Asp Ala Ala Thr Asp Lys Val Ala Thr Ala  
450 455 460

Ser Glu Gln Thr Gly Trp Lys Tyr Thr Phe Thr Asn Leu Pro Lys Tyr  
465 470 475 480

Lys Asp Gly Lys Gln Ile Thr Tyr Thr Ile Gln Glu Asp Pro Val Ala  
485 490 495

Asp Tyr Thr Thr Ile Gln Gly Phe Asp Ile Thr Asn His His Glu  
500 505 510

Val Ala Leu Thr Ser Leu Lys Val Ile Lys Val Trp Asn Asp Lys Asp  
515 520 525

Asp Tyr Tyr His Lys Arg Pro Lys Glu Ile Thr Ile Leu Leu Lys Ala  
530 535 540

Asp Gly Lys Val Ile Arg Glu His Gln Met Thr Pro Asp Gln Gln Gly  
545 550 555 560

Lys Trp Glu Tyr Thr Phe Asp Gln Leu Pro Val Tyr Gln Thr Gly Lys  
565 570 575

Lys Ile Ser Tyr Ser Ile Glu Glu Lys Gln Val Ala Gly Tyr Gln Ala  
580 585 590

Pro Val Tyr Glu Val Asp Glu Gly Leu Lys Gln Val Thr Val Thr Asn  
595 600 605

Thr Leu Asn Pro Ser Tyr Lys Leu Pro Asp Thr Gly Gly Gln Gly Val  
610 615 620

Lys Trp Tyr Leu Leu Ile Gly Gly Phe Ile Ile Val Ala Ile Leu  
625 630 635 640

Val Leu Ile Ser Leu Tyr Gln Lys His Lys Arg His Asn Met Ser Lys  
645 650 655

Pro

110087400US.ST25

&lt;211&gt; 34

&lt;212&gt; DNA

&lt;213&gt; Oligonucleotide primer

&lt;400&gt; 5

ttttctcgag ctacggtaga gctgataaaaa tctc

34

&lt;210&gt; 6

&lt;211&gt; 32

&lt;212&gt; DNA

&lt;213&gt; Oligonucleotide primer

&lt;400&gt; 6

tcagccatgg ctctagatgc tacaacggtg tt

32

&lt;210&gt; 7

&lt;211&gt; 600

&lt;212&gt; DNA

&lt;213&gt; Streptococcus equi

&lt;400&gt; 7

aaataatttt gtttaacttt aagaaggaga tataaccatg gctctagatg ctacaacggt

60

gttagagcct acaacagcct tcatttagaga agctgttagg gaaatcaatc agctgagtga

120

tgactacgct gacaatcaag agttcaggc tggcttgct aatgctggag ttgaggcact

180

tgctgcagat actgttgatc aggctaaagc agctcttgac aaagcaaagg cagctgtgc

240

tgggtttcag cttgtatgaag caagacgtga ggcttacaga acaatcaatg ccttaagtga

300

tcagcacaaa agcgatcaaa aggttcagct agctcttagtt gctgcagcag ctaaggtggc

360

agatgctgct tcagttgatc aagtgaatgc agccattaat gatgctcata cagctattgc

420

ggacattaca ggagcagcct tggggaggc taaagaagct gctatcaatg aactaaagca

480

gtatggcatt agtgattact atgtgacctt aatcaacaaa gccaaaactg ttgaaggtgt

540

caatgcgctt aaggcaaaga ttttatcagc tctaccgtag ctcgagcccg ggtgcttgc

600

&lt;210&gt; 8

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Oligonucleotide primer

## 110087400US.ST25

<400> 8  
ggtcccatgg caactccgaa tttagaagga 30

<210> 9

<211> 29

<212> DNA

<213> oligonucleotide primer

<400> 9  
cagactcgag gtcgggattg taagaatag 29

<210> 10

<211> 125

<212> PRT

<213> *Streptococcus equi*

<400> 10

Met Ala Thr Pro Asn Leu Glu Gly Asn Asn Arg Lys Asn Pro Leu Lys  
1 5 10 15

Gly Leu Asp Gly Glu Asn Lys Pro Lys Glu Asp Leu Asp Gly Lys Gly  
20 25 30

Leu Ser Gly Glu Asn Asp Glu Ser Pro Lys Leu Lys Asp Glu His Pro  
35 40 45

Tyr Asn His Gly Arg Arg Asp Gly Tyr Arg Val Gly Tyr Glu Asp Gly  
50 55 60

Tyr Gly Gly Lys Lys His Lys Gly Asp Tyr Pro Lys Arg Phe Asp Glu  
65 70 75 80

Ser Ser Pro Lys Glu Tyr Asn Asp Tyr Ser Gln Gly Tyr Asn Asp Asn  
85 90 95

Tyr Gly Asn Gly Tyr Leu Asp Gly Leu Ala Asp Arg Gly Gly Lys Arg  
100 105 110

Gly Tyr Gly Tyr Ser Tyr Asn Pro Asp Leu Glu Pro Gly  
115 120 125

<210> 11

<211> 34

<212> DNA

110087400US.ST25

&lt;213&gt; Oligonucleotide primer

<400> 11  
accatggcta gcgcagagca gctttattat gggt

34

&lt;210&gt; 12

&lt;211&gt; 32

&lt;212&gt; DNA

&lt;213&gt; Oligonucleotide primer

<400> 12  
atacccgga tatttcgg tactaccata gt

32

&lt;210&gt; 13

&lt;211&gt; 310

&lt;212&gt; PRT

&lt;213&gt; Streptococcus equi

&lt;400&gt; 13

Met Ala Ser Ala Glu Gln Leu Tyr Tyr Gly Trp Asn Asp Gly Thr Arg  
1 5 10 15Gln Ser Ser Pro Tyr Phe Leu Tyr Val Ser Pro Lys Asn Ala Pro Lys  
20 25 30Arg Glu Leu Lys Asp Glu Tyr Val Val Tyr Cys Phe Asn Lys Lys Leu  
35 40 45Tyr Trp Pro Asp Gln Trp Glu Ser Ile Tyr Ser Asn Phe Asn Asp Ile  
50 55 60Arg Ser Pro Tyr Asn Asp Leu Pro Val Tyr Glu Lys Lys Leu Gly Tyr  
65 70 75 80Asp Gly Ile Phe Lys Gln Tyr Ala Pro Asp Tyr Lys Lys Asp Ile Ser  
85 90 95Asp Ile Ala Ser Ala Leu Val Ala Val Leu Ser Asn Gly Tyr Pro Thr  
100 105 110Asn Lys Ser Gln Leu Ser Thr Ser Tyr His Leu Asn Asn Asp Ser Ser  
115 120 125Arg Lys Val Thr Gln Leu Ala Ile Trp Tyr Phe Ser Asp Ser Leu Thr  
130 135 140

110087400US.ST25

Lys Glu Tyr Leu Lys Asp Thr Gly Gly Tyr Asn Leu Asn Asp Met Glu  
145 150 155 160

Lys Lys Ala Leu Asp Phe Leu Ile Ser Lys Gly Glu Asp Ser Lys Leu  
165 170 175

Lys Ser Glu Gln Ser Asn Tyr Ser Leu Asp Ile Tyr Val Tyr Gln Ser  
180 185 190

Gly Gly His Asp His Met Lys Asp Tyr Gln Asn Leu Leu Gly Ser Thr  
195 200 205

Leu Ile Pro Lys Glu Pro Leu Lys Pro Gln Leu Gly Gly Phe Ser Gly  
210 215 220

His Asn Gly Asn Gly Leu Ser Gly Leu Glu Gly Gly Ser Ser Gly Ser  
225 230 235 240

Gln Glu Thr Asn Glu Asp Gly Lys Lys Gly Leu Ile Gly Phe His Gly  
245 250 255

Gly Leu Ser Gly Ser Glu Gly Lys Arg Asp Pro Leu Pro Gly Leu Lys  
260 265 270

Gly Glu Ala Gly Ala Pro Asp Thr Pro Gln Lys Pro Asn Asp Pro Leu  
275 280 285

Gln Gly Leu Glu Gly Gly Asn Ser Pro Ile Val Glu Gln Asn Tyr Gly  
290 295 300

Ser Thr Glu Gly Tyr Gly  
305 310

<210> 14

<211> 5

<212> PRT

<213> C-terminal motif

<400> 14

Leu Pro Asp Thr Gly  
1 5

<210> 15

<211> 1971

<212> DNA

110087400US.ST25

&lt;213&gt; Streptococcus equi

<400>	15					
ttgaaaacaac	tgacaaagat	cgttagtgtg	gtcttggc	tggctttac	ccttagtgct	60
agcctgcaca	aggttcggc	aactaatctt	agtgacaaca	tcacatcatt	gacggttgct	120
tcttcatcac	tccgagatgg	agagagaacg	acggtaaagg	ttgcgttga	tgacaaaaaa	180
cagaaaatca	aggcagggga	tacgatagag	gtcacctggc	ctacaagtgg	taatgtctac	240
attcagggct	ttaataaaac	cataccgctt	aatatttagag	gggttagatgt	tggtaccttg	300
gaggtcacgc	tagacaaggc	tgtttcaca	ttcaatcaa	atattgaaac	aatgcatgat	360
gtctctggtt	ggggagagtt	tgatattact	gttagaaatg	tgacacaaac	caccgctgaa	420
acatcaggaa	cgaccacagt	aaaggttaggc	aatcgcactg	ctactatcac	tgttactaag	480
cctgaggcag	gcactggcac	cagctcattt	tattataaga	ctggtgat	gcagcccaat	540
gatactgagc	gtgtgagatg	gttccctgctg	attaacaaca	acaaggaatg	ggtggccaat	600
actgttacag	tcgaagacga	tattcaaggt	ggtcaaacct	tggatatgag	cagcttgac	660
atcaccgtat	ctggttatcg	taacgagcgc	ttcggtgggg	aaaacgctct	gacagagttt	720
catacaacat	ttccaaattc	tgtcattacg	gcaacagata	atcacattag	tgtcggtta	780
gatcaatatg	atgcctcaca	aaacactgtc	aacattgctt	ataagacaaa	gataacggac	840
tttgaccaaa	aagaatttgc	caacaacagt	aaaatctgg	accagatttt	atacaaggat	900
caggtatcgg	gtcaagagtc	aaaccaccaa	gtagccaata	tcaatgctaa	cggcggggtt	960
gatggcagtc	gctataccag	cttactgtc	aagaaaattt	ggaatgacaa	ggaaaatcaa	1020
gacggtaagc	gtccaaagac	tattactgtt	cagcttacg	ccaatgatca	gaaagttaat	1080
gataagacca	ttgaatttgc	tgataactaat	agctggcaag	caagtttgg	taagctggat	1140
aagtatgaca	gtcagaacca	aaaaattacc	tacagtgtca	aggaagtgtat	ggttccctgtt	1200
ggctaccaat	cgcagggtga	gggggatagt	ggagtaggat	ttaccattac	caacacctat	1260
acaccagagg	tcattagcat	taccggcaa	aaaacttggg	acgacaggga	aaaccaagac	1320
ggtaaacgtc	ctaaggagat	tacggttcgt	ttattggcaa	atgacgctgc	aactgacaag	1380
gtagcaactg	tttcagagca	aaccggctgg	aagtatacat	ttaccaatct	accgaaatac	1440
aaagatggta	aacagatcac	ctacacgatc	caagaggacc	ctgtggcaga	ttacaccaca	1500
accattcagg	gatttgcata	taccaatcat	catgaggtag	ccttgaccag	cctaaaggtc	1560
atcaagggtt	ggaatgataa	ggacgattat	taccataaac	gtcccaagga	gattaccatt	1620
ttgctaaagg	cagatggcaa	ggtagttcgt	gaacatcaga	tgacaccgga	tcagcaagga	1680
aaatgggaat	acaccttgc	ccagctgccc	gtctatcaga	caggcaagaa	aatcagctac	1740
agcattgagg	aaaaacaggt	tgctggctat	caagcccctg	tctatgaggt	tgatgaaggc	1800
ttgaaggcagg	tcactgtaac	caacaccctt	aacccaagct	acaagctgcc	tgacaccgga	1860
ggacaaggag	tgaaatggta	cctgttaatc	ggtggcggtt	ttatcatcgt	cgcaatcctt	1920

110087400US.ST25

gtactgatca gccttatca aaaacacaag cgccataaca tgtcaaaacc a 1971

<210> 16

<211> 32

<212> DNA

<213> Oligonucleotide primer

<400> 16  
catgccatgg caactaatct tagtgacaac at 32

<210> 17

<211> 32

<212> DNA

<213> Oligonucleotide primer

<400> 17  
ccgctcgagc ttgttagcttg ggttaagggt gt 32

<210> 18

<211> 32

<212> DNA

<213> Oligonucleotide primer

<400> 18  
ccgctcgaga aagctggtat agcgactgcc at 32

<210> 19

<211> 1782

<212> DNA

<213> *Streptococcus equi*

<400> 19  
atggcaacta atcttagtga caacatcaca tcattgacgg ttgcttcttc atcactccga 60  
gatggagaga gaacgacggt aaagggttgcg tttgatgaca aaaaacagaa aatcaaggca 120  
ggggatacga tagaggtcac ctggcctaca agtggtaatg tctacattca gggctttaat 180  
aaaaccatac cgcttaatat tagagggta gatgttgta ccttggaggt cacgctagac 240  
aaggctgttt tcacattcaa tcaaaatatt gaaacaatgc atgatgtctc tggttgggga 300  
gagtttgata ttactgttag aatgtgaca caaaccaccg ctgaaacatc aggaacgacc 360

## 110087400US.ST25

acagtaaagg taggcaatcg cactgctact atcactgtta ctaagcctga ggcaggcact	420
ggtaccagct cattttatta taagactggt gatattcagc ccaatgatac tgagcgtgtg	480
agatggttcc tgctgattaa caacaacaag gaatgggtgg ccaatactgt tacagtcga	540
gacgatattc aaggtggtca aacccttggat atgagcagct ttgacatcac cgtatctgg	600
tatcgtaacg agcgcttcgt tggggaaaac gctctgacag agtttcatac aacatttcca	660
aattctgtca ttacggcaac agataatcac attagtgtgc ggtagatca atatgatgcc	720
tcacaaaaca ctgtcaacat tgcttataag acaaagataa cggactttga ccaaaaagaa	780
tttgccaaaca acagtaaaat ctggtaccag attttatatac aggatcaggt atcgggtcaa	840
gagtcaaacc accaagttagc caatatcaat gctaacggcg gggttgatgg cagtcgtat	900
accagcttta ctgtcaagaa aatttggaat gacaaggaaa atcaagacgg taagcgtcca	960
aagactatta ctgttcagct ttacgccaat gatcagaaag ttaatgataa gaccattgaa	1020
ttgagtgata ctaatagctg gcaagcaagt tttggtaagc tggataagta tgacagtcag	1080
aaccaaaaaa ttacctacag tgtcaaggaa gtgatggttc ctgttggcta ccaatcgca	1140
gttgggggg atagtggagt aggatttacc attaccaaca cctatacacc agaggtcatt	1200
agcattaccg gtcaaaaaac ttgggacgac agggaaaacc aagacggtaa acgtcctaag	1260
gagattacgg ttcgtttatt ggcaaatgac gctgcaactg acaaggtagc aactgcttca	1320
gagcaaaccg gctggaagta tacatttacc aatctaccga aatacaaaga tggtaaacag	1380
atcacctaca cgatccaaga ggaccctgtg gcagattaca ccacaaccat tcagggattt	1440
gatattacca atcatcatga ggtgccttg accagcctaa aggtcatcaa ggtttggaat	1500
gataaggacg attattacca taaacgtccc aaggagatta ccattttgct aaaggcagat	1560
ggcaaggtga ttcgtgaaca tcagatgaca ccggatcagc aaggaaaatg ggaatacacc	1620
tttgaccagc tgccggctta tcagggcaggc aagaaaatca gctacagcat tgaggaaaaa	1680
caggttgctg gctatcaagc ccctgtctat gaggttgatg aaggcttgaa gcaggtcact	1740
gtaaccaaca cccttaaccc aagctacaag ctcgagcccg gg	1782

&lt;210&gt; 20

&lt;211&gt; 594

&lt;212&gt; PRT

<213> *Streptococcus equi*

&lt;400&gt; 20

Met Ala Thr Asn Leu Ser Asp Asn Ile Thr Ser Leu Thr Val Ala Ser	
1	5
10	15

Ser Ser Leu Arg Asp Gly Glu Arg Thr Thr Val Lys Val Ala Phe Asp	
20	25
30	

110087400US.ST25

Asp Lys Lys Gln Lys Ile Lys Ala Gly Asp Thr Ile Glu Val Thr Trp  
35 40 45

Pro Thr Ser Gly Asn Val Tyr Ile Gln Gly Phe Asn Lys Thr Ile Pro  
50 55 60

Leu Asn Ile Arg Gly Val Asp Val Gly Thr Leu Glu Val Thr Leu Asp  
65 70 75 80

Lys Ala Val Phe Thr Phe Asn Gln Asn Ile Glu Thr Met His Asp Val  
85 90 95

Ser Gly Trp Gly Glu Phe Asp Ile Thr Val Arg Asn Val Thr Gln Thr  
100 105 110

Thr Ala Glu Thr Ser Gly Thr Thr Val Lys Val Gly Asn Arg Thr  
115 120 125

Ala Thr Ile Thr Val Thr Lys Pro Glu Ala Gly Thr Gly Thr Ser Ser  
130 135 140

Phe Tyr Tyr Lys Thr Gly Asp Ile Gln Pro Asn Asp Thr Glu Arg Val  
145 150 155 160

Arg Trp Phe Leu Leu Ile Asn Asn Asn Lys Glu Trp Val Ala Asn Thr  
165 170 175

Val Thr Val Glu Asp Asp Ile Gln Gly Gln Thr Leu Asp Met Ser  
180 185 190

Ser Phe Asp Ile Thr Val Ser Gly Tyr Arg Asn Glu Arg Phe Val Gly  
195 200 205

Glu Asn Ala Leu Thr Glu Phe His Thr Thr Phe Pro Asn Ser Val Ile  
210 215 220

Thr Ala Thr Asp Asn His Ile Ser Val Arg Leu Asp Gln Tyr Asp Ala  
225 230 235 240

Ser Gln Asn Thr Val Asn Ile Ala Tyr Lys Thr Lys Ile Thr Asp Phe  
245 250 255

Asp Gln Lys Glu Phe Ala Asn Asn Ser Lys Ile Trp Tyr Gln Ile Leu  
260 265 270

Tyr Lys Asp Gln Val Ser Gly Gln Glu Ser Asn His Gln Val Ala Asn  
275 280 285

Ile Asn Ala Asn Gly Gly Val Asp Gly Ser Arg Tyr Thr Ser Phe Thr  
290 295 300

110087400US.ST25

Val Lys Lys Ile Trp Asn Asp Lys Glu Asn Gln Asp Gly Lys Arg Pro  
305 310 315 320

Lys Thr Ile Thr Val Gln Leu Tyr Ala Asn Asp Gln Lys Val Asn Asp  
325 330 335

Lys Thr Ile Glu Leu Ser Asp Thr Asn Ser Trp Gln Ala Ser Phe Gly  
340 345 350

Lys Leu Asp Lys Tyr Asp Ser Gln Asn Gln Lys Ile Thr Tyr Ser Val  
355 360 365

Lys Glu Val Met Val Pro Val Gly Tyr Gln Ser Gln Val Glu Gly Asp  
370 375 380

Ser Gly Val Gly Phe Thr Ile Thr Asn Thr Tyr Thr Pro Glu Val Ile  
385 390 395 400

Ser Ile Thr Gly Gln Lys Thr Trp Asp Asp Arg Glu Asn Gln Asp Gly  
405 410 415

Lys Arg Pro Lys Glu Ile Thr Val Arg Leu Leu Ala Asn Asp Ala Ala  
420 425 430

Thr Asp Lys Val Ala Thr Ala Ser Glu Gln Thr Gly Trp Lys Tyr Thr  
435 440 445

Phe Thr Asn Leu Pro Lys Tyr Lys Asp Gly Lys Gln Ile Thr Tyr Thr  
450 455 460

Ile Gln Glu Asp Pro Val Ala Asp Tyr Thr Thr Ile Gln Gly Phe  
465 470 475 480

Asp Ile Thr Asn His His Glu Val Ala Leu Thr Ser Leu Lys Val Ile  
485 490 495

Lys Val Trp Asn Asp Lys Asp Asp Tyr Tyr His Lys Arg Pro Lys Glu  
500 505 510

Ile Thr Ile Leu Leu Lys Ala Asp Gly Lys Val Ile Arg Glu His Gln  
515 520 525

Met Thr Pro Asp Gln Gln Gly Lys Trp Glu Tyr Thr Phe Asp Gln Leu  
530 535 540

Pro Val Tyr Gln Ala Gly Lys Lys Ile Ser Tyr Ser Ile Glu Glu Lys  
545 550 555 560

Gln Val Ala Gly Tyr Gln Ala Pro Val Tyr Glu Val Asp Glu Gly Leu  
565 570 575

110087400US.ST25

Lys Gln Val Thr Val Thr Asn Thr Leu Asn Pro Ser Tyr Lys Leu Glu  
 580 585 590

Pro Gly

<210> 21

<211> 921

<212> DNA

<213> *Streptococcus equi*

<400> 21  
 atggcaacta atcttagtga caacatcaca tcattgacgg ttgcttcttc atcactccga 60  
 gatggagaga gaacgacggt aaagggtgcg tttgatgaca aaaaacagaa aatcaaggca 120  
 ggggatacga tagaggtcac ctggcctaca agtggtaatg tctacattca gggctttaat 180  
 aaaaccatac cgcttaatat tagagggta gatgttgta cttggaggt cacgctagac 240  
 aaggctgttt tcacattcaa tcaaaatatt gaaacaatgc atgatgtctc tgggtgggaa 300  
 gagtttgata ttactgttag aaatgtgaca caaaccaccg ctgaaacatc aggaacgacc 360  
 acagtaaagg taggcaatcg cactgctact atcactgtta ctaaggctga ggcaggcact 420  
 ggtaccagct cattttatta taagactggt gatatgcagc ccaatgatac tgagcgtgtg 480  
 agatggttcc tgctgattaa caacaacaag gaatgggtgg ccaatactgt tacagtcga 540  
 gacgatattc aaggtggtca aaccttggat atgagcagct ttgacatcac cgtatctgg 600  
 tatacgtaacg agcgcttcgt tggggaaaac gctctgacag agtttcatac aacatttcca 660  
 aattctgtca ttacggcaac agataatcac attagtgtgc ggttagatca atatgatgcc 720  
 tcacaaaaca ctgtcaacat tgcttataag acaaagataa cggactttga ccaaaaagaa 780  
 tttgccaaca acagtaaaat ctggtaccag attttataca aggatcaggt atcgggtcaa 840  
 gagtc当地 accaagttagc caatatcaat gctaacggcg ggggtgatgg cagtcgtat 900  
 accagcttcc tcgagcccg g 921

<210> 22

<211> 307

<212> PRT

<213> *Streptococcus equi*

<400> 22

Met Ala Thr Asn Leu Ser Asp Asn Ile Thr Ser Leu Thr Val Ala Ser  
 1 5 10 15

110087400US.ST25

Ser Ser Leu Arg Asp Gly Glu Arg Thr Thr Val Lys Val Ala Phe Asp  
20 25 30

Asp Lys Lys Gln Lys Ile Lys Ala Gly Asp Thr Ile Glu Val Thr Trp  
35 40 45

Pro Thr Ser Gly Asn Val Tyr Ile Gln Gly Phe Asn Lys Thr Ile Pro  
50 55 60

Leu Asn Ile Arg Gly Val Asp Val Gly Thr Leu Glu Val Thr Leu Asp  
65 70 75 80

Lys Ala Val Phe Thr Phe Asn Gln Asn Ile Glu Thr Met His Asp Val  
85 90 95

Ser Gly Trp Gly Glu Phe Asp Ile Thr Val Arg Asn Val Thr Gln Thr  
100 105 110

Thr Ala Glu Thr Ser Gly Thr Thr Val Lys Val Gly Asn Arg Thr  
115 120 125

Ala Thr Ile Thr Val Thr Lys Pro Glu Ala Gly Thr Gly Thr Ser Ser  
130 135 140

Phe Tyr Tyr Lys Thr Gly Asp Met Gln Pro Asn Asp Thr Glu Arg Val  
145 150 155 160

Arg Trp Phe Leu Leu Ile Asn Asn Asn Lys Glu Trp Val Ala Asn Thr  
165 170 175

Val Thr Val Glu Asp Asp Ile Gln Gly Gln Thr Leu Asp Met Ser  
180 185 190

Ser Phe Asp Ile Thr Val Ser Gly Tyr Arg Asn Glu Arg Phe Val Gly  
195 200 205

Glu Asn Ala Leu Thr Glu Phe His Thr Thr Phe Pro Asn Ser Val Ile  
210 215 220

Thr Ala Thr Asp Asn His Ile Ser Val Arg Leu Asp Gln Tyr Asp Ala  
225 230 235 240

Ser Gln Asn Thr Val Asn Ile Ala Tyr Lys Thr Lys Ile Thr Asp Phe  
245 250 255

Asp Gln Lys Glu Phe Ala Asn Asn Ser Lys Ile Trp Tyr Gln Ile Leu  
260 265 270

Tyr Lys Asp Gln Val Ser Gly Gln Glu Ser Asn His Gln Val Ala Asn  
275 280 285

110087400US.ST25

Ile Asn Ala Asn Gly Gly Val Asp Gly Ser Arg Tyr Thr Ser Phe Leu  
290 295 300

Glu Pro Gly  
305

<210> 23

<211> 302

<212> PRT

<213> Streptococcus equi

<400> 23

Met Thr Asn Lys Thr Lys Arg Thr Gly Leu Val Arg Lys Tyr Gly Ala  
1 5 10 15

Cys Ser Ala Ala Ile Ala Leu Ala Ala Leu Ala Ser Leu Gly Ala Gly  
20 25 30

Lys Ala Val Lys Ala Asp Gln Pro Ala Ala Leu Lys Tyr Pro Glu Pro  
35 40 45

Arg Asp Tyr Phe Leu His Thr Arg Glu Gly Asp Val Ile Tyr Asp Glu  
50 55 60

Asp Ile Lys Arg Tyr Phe Glu Asp Leu Glu Ala Tyr Leu Thr Ala Arg  
65 70 75 80

Leu Gly Gly Ile Asp Lys Lys Val Glu Glu Ala Ala Gln Lys Pro Gly  
85 90 95

Ile Pro Gly Pro Thr Gly Pro Gln Gly Pro Lys Gly Asp Lys Gly Asp  
100 105 110

Pro Gly Ala Pro Gly Glu Arg Gly Pro Ala Gly Pro Lys Gly Asp Thr  
115 120 125

Gly Glu Ala Gly Pro Arg Gly Glu Gln Gly Pro Ala Gly Gln Ala Gly  
130 135 140

Glu Arg Gly Pro Lys Gly Asp Pro Gly Ala Pro Gly Pro Lys Gly Glu  
145 150 155 160

Lys Gly Asp Thr Gly Ala Val Gly Pro Lys Gly Glu Lys Gly Asp Thr  
165 170 175

Gly Ala Thr Gly Pro Lys Gly Asp Lys Gly Glu Arg Gly Glu Lys Gly  
180 185 190

110087400US.ST25

Glu Gln Gly Gln Arg Gly Glu Lys Gly Glu Gln Gly Gln Arg Gly Glu  
 195 200 205

Lys Gly Glu Gln Lys Pro Lys Gly Asp Gln Gly Lys Asp Thr Lys Pro  
 210 215 220

Ser Ala Pro Lys Ala Pro Glu Lys Ala Pro Ala Pro Lys Ala Pro Lys  
 225 230 235 240

Ala Ser Glu Gln Ser Ser Asn Pro Lys Ala Pro Ala Pro Lys Ser Ala  
 245 250 255

Pro Ser Lys Ser Ala Ala Pro Thr Gly Gln Lys Ala Ala Leu Pro Ala  
 260 265 270

Thr Gly Glu Ile Asn His Pro Phe Phe Thr Leu Ala Ala Leu Ser Val  
 275 280 285

Ile Ala Ser Val Gly Val Leu Thr Leu Lys Gly Lys Lys Asp  
 290 295 300

<210> 24

<211> 909

<212> DNA

<213> *Streptococcus equi*

<400> 24		
atgacaaaca aaacaaagcg tacaggattg gtacgcaagt acgggtgcctg ctcagcagct	60	
atcgcccttag cagctcttgc aagcctggga gcaggtaaag cagtaaaggc agaccagcca	120	
gcagcactaa aatatccaga acctagagac tattttcttc atactcgtga aggtgatgtt	180	
atttatgtg aggatataaa aagatatttt gaggatttag aagcctattt aacagctaga	240	
cttgggtggga ttgataaaaa agtagaagaa gctgccccaa agccaggtat tccaggtcct	300	
actggccctc aaggtcctaa gggagacaaa ggagatccag gtgccccctgg tgagcgcgg	360	
ccagctggac caaaggcgca tacgggcgaa gcccggacaa gaggtgagca aggcccagcc	420	
ggacaagctg gagaacgtgg accaaaagga gatccaggtg ctccaggtcc taaaggtgaa	480	
aagggtgata ctggtgcagt tggtcctaaa ggtgaaaaag gtgataaccgg agcaaccgga	540	
ccaaaggggag acaaggggcgaa acgcggtgaa aaaggcgagc aaggccaaacg tggcgaaaaaa	600	
ggcgagcaag gccaacgcgg tgaaaaaggc gagcaaaaac caaagggtga tcaaggaaaa	660	
gatacaaaaac catcagctcc aaaagcacct gaaaaggctc ctgcacccaa agctccaaag	720	
gcttcagagc agtcatctaa tcctaaagca ccagctccctaa agtcagcacc aagcaaatca	780	
gcggcaccaa caggtcaaaa agcagcccta ccagcaacag gggaaatcaa ccaccatc	840	

## 110087400US.ST25

ttcacccttg cagctcttag tgtcatcgct agcgtaggcg tcctaactct aaaaggaaaa 900  
aaagactaa 909

<210> 25

<211> 33

<212> DNA

<213> Oligonucleotide primer

<400> 25  
catgccccatgg accagccagc agcactaaaa tat 33

<210> 26

<211> 31

<212> DNA

<213> Oligonucleotide primer

<400> 26  
ccgctcgagg gctgctttt gacctgttgg t 31

<210> 27

<211> 237

<212> PRT

<213> Streptococcus equi

<400> 27

Met Asp Gln Pro Ala Ala Leu Lys Tyr Pro Glu Pro Arg Asp Tyr Phe  
1 5 10 15

Leu His Thr Arg Glu Gly Asp Val Ile Tyr Asp Glu Asp Ile Lys Arg  
20 25 30

Tyr Phe Glu Asp Leu Glu Ala Tyr Leu Thr Ala Arg Leu Gly Gly Ile  
35 40 45

Asp Lys Lys Val Glu Glu Ala Ala Gln Lys Pro Gly Ile Pro Gly Pro  
50 55 60

Thr Gly Pro Gln Gly Pro Lys Gly Asp Lys Gly Asp Pro Gly Ala Pro  
65 70 75 80

Gly Glu Arg Gly Pro Ala Gly Pro Lys Gly Asp Thr Gly Glu Ala Gly  
85 90 95

110087400US.ST25

Pro Arg Gly Glu Gln Gly Pro Ala Gly Gln Ala Gly Glu Arg Gly Pro  
100 105 110

Lys Gly Asp Pro Gly Ala Pro Gly Pro Lys Gly Glu Lys Gly Asp Thr  
115 120 125

Gly Ala Val Gly Pro Lys Gly Glu Lys Gly Asp Thr Gly Ala Thr Gly  
130 135 140

Pro Lys Gly Asp Lys Gly Glu Arg Gly Glu Lys Gly Glu Gln Gly Gln  
145 150 155 160

Arg Gly Glu Lys Gly Glu Gln Gly Gln Arg Gly Glu Lys Gly Glu Gln  
165 170 175

Lys Pro Lys Gly Asp Gln Gly Lys Asp Thr Lys Pro Ser Ala Pro Lys  
180 185 190

Ala Pro Glu Lys Ala Pro Ala Pro Lys Ala Pro Lys Ala Ser Glu Gln  
195 200 205

Ser Ser Asn Pro Lys Ala Pro Ala Pro Lys Ser Ala Pro Ser Lys Ser  
210 215 220

Ala Ala Pro Thr Gly Gln Lys Ala Ala Leu Glu Pro Gly  
225 230 235

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/SE 2003/001587

## A. CLASSIFICATION OF SUBJECT MATTER

IPC7: A61K 39/09, C07K 14/315, C07K 16/12

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC7: C07K, A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

SE,DK,FI,NO classes as above

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

EPO-INTERNAL, WPI-DATA, PAJ, BIOSIS, MEDLINE, CHEM. ABS DATA

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	FEMS Microbiology Letters, Volume 222, 2003, Jonas Lannergård et al: "CNE, a collagen-binding protein of Streptococcus equi", page 69 - page 74, abstract; page 70, column 2, paragraph 1; page 73, column 2, paragraph 2	12,19
P,Y	--	3,5,7-19
X	Hans Lindmark, "Acta Universitatis Agriculturae Sueciae, AGRARIA 139, Characterization of Adhesive Extracellular Proteins from Streptococcus equi", 1999, Doctoral thesis Swedish University of Agricultural Sciences, Uppsala, page 1 - page 49, abstract; page 25 - page 35	1-2,4-19
Y	--	2-5,7-19

 Further documents are listed in the continuation of Box C. See patent family annex.

\* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or conflict underlying the invention

"X" document of particular relevance: the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance: the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&amp;" document member of the same patent family

Date of the actual completion of the international search

12 January 2004

Date of mailing of the international search report

20-01-2004

Name and mailing address of the ISA/  
Swedish Patent Office  
Box 5055, S-102 42 STOCKHOLM  
Facsimile No. + 46 8 666 02 86

Authorized officer

Terese Persson/EÖ  
Telephone No. + 46 8 782 25 00

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/SE 2003/001587

## C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	Hans Lindmark et al, "Acta Universitatis Agricuture Sueciae, AGRARIA 139, Characterization of Adhesive Extracellular Proteins from Streptococcus equi", 1999, The NH2-terminal half of protein FNZ binds fibronectin and is secreted in S. equi subsp. equi but not in subsp. zooepidemicus, Swedish University of Agricultural Sciences, Uppsala, page1 - page 13, abstract; page 93, column 2, paragraph 2; page 96	1-2,4-19
Y	--	2-5,7-19
X	Research in Veterinary Science, Volume 66, 1999, Hans Lindmark et al: "Pulsed-field gel electrophoresis and distribution of the genes zag and fnz in isolates of Streptococcus equi", page 93 - page 99, abstract; page 93, column 2, paragraph 2; page 96	1-2,4-19
Y	--	2,5,7-19
X	Infection and Immunity, Volume 67, no. 5, May 1999, Hans Lindmark et al: "SFS, a Novel Fibronectin-Binding Protein from Streptococcus equi, Inhibits the Binding between Fibronectin and Collagen", page 2383 - page 2388, abstract; page 2384, column 1, paragraph 5	12,19
Y	--	2,5,7-19
X	Infection and Immunity, Volume 69, no. 5, May 2001, Hans Lindmark et al: "Comparison of the Fibronectin-Binding Protein FNE from Streptococcus equi Subspecies equi with FNZ from S. equi Subspecies zooepidemicus Reveals a Major and Conserved Difference", page 3159 - page 3163, abstract; page 3160, column 2, paragraph 3	12,19
Y	--	2,4,5,7-19

## INTERNATIONAL SEARCH REPORT

International application No.

PCT/SE 2003/001587

## C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 0037496 A1 (GUSS, BENGT ET AL), 29 June 2000 (29.06.2000), claims	12,19
Y	--	2,5,7-19
X	WO 9507296 A1 (GUSS, BENGT ET AL), 16 March 1995 (16.03.1995), abstract; claims	1,6,8-9,11
Y	-- -----	2,3,7-19

# INTERNATIONAL SEARCH REPORT

International application No.  
**PCT/SE03/01587**

## Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1.  Claims Nos.: **16-18**  
because they relate to subject matter not required to be searched by this Authority, namely:  
**see next sheet\***
2.  Claims Nos.: **20 (partially)**  
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:  
**see next sheet\*\***
3.  Claims Nos.:  
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

## Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1.  As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2.  As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3.  As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4.  No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

### Remark on Protest

The additional search fees were accompanied by the applicant's protest.  
 No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International application No.  
PCT/SE03/01587

\*

Claims 16-18 relate to methods of treatment of the human or animal body by surgery or by therapy or diagnostic methods practised on the human or animal body (PCT Rule 39.1(iv)). Nevertheless, a search has been executed for these claims. The search has been based on the alleged effects of the compounds or compositions.

\*\*

Present claim 20 relates to an extremely large number of possible protein fragments due to the expression "fragment thereof". Support within the meaning of Article 6 PCT and disclosure within the meaning of Article 5 PCT is to be found, however, for only a very small proportion of the protein fragments. In the present case, the claim so lacks support, and the application so lacks disclosure, that a meaningful search over the whole of the claimed scope is impossible.

Consequently, the search has been carried out for those parts of the claims which appear to be supported and disclosed, namely those parts related to the protein SclC and the fragment SclC1, as well as general aspects thereof.

The applicant's attention is drawn to the fact that claims, or parts of claims, relating to inventions in respect of which no international search report has been established will not be, the subject of an international preliminary examination (Rule 66.1(e) PCT). This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure.

INTERNATIONAL SEARCH REPORT  
Information on patent family members

01/12/2003

International application No.  
PCT/SE 2003/001587

WO	0037496	A1	29/06/2000	AU	3094700	A	12/07/2000
				CA	2355831	A	29/06/2000
				EP	1151003	A	07/11/2001
				SE	9804491	D	00/00/0000
				US	2003165527	A	04/09/2003

WO	9507296	A1	16/03/1995	AT	194360	T	15/07/2000
				AU	7628194	A	27/03/1995
				CA	2176230	A	16/03/1995
				DE	69425149	D,T	15/03/2001
				EP	0724596	A,B	07/08/1996
				ES	2149884	T	16/11/2000
				SE	9302855	D	00/00/0000
				US	6100055	A	08/08/2000